

40220" E140T660

Q9925764.Seq

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Q9925765 . Seq  
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Q9925767. Seq

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09925768. Seq

Q9925/68.Seq

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Q9925768.Seq  
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Test 466

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Q9925779.1 Seq

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#44540 54407650

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4640" E460T.660

Q9925796.Seq

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For 240" 2490T660

Q9925797.Seq

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Q9925798 . Seq

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1990	1991	1992	1993	1994	1995	1996	1997	1998	1999	2000	2001	2002	2003	2004	2005	2006	2007	2008	2009	2010	2011	2012	2013	2014	2015	2016	2017	2018	2019	2020	2021	2022	2023	2024	2025	2026	2027	2028	2029	2030	2031	2032	2033	2034	2035	2036	2037	2038	2039	2040	2041	2042	2043	2044	2045	2046	2047	2048	2049	2050	2051	2052	2053	2054	2055	2056	2057	2058	2059	2060	2061	2062	2063	2064	2065	2066	2067	2068	2069	2070	2071	2072	2073	2074	2075	2076	2077	2078	2079	2080	2081	2082	2083	2084	2085	2086	2087	2088	2089	2090	2091	2092	2093	2094	2095	2096	2097	2098	2099	2100	

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442240 24647663

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Q9925891.Seq

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CAGTACATTTCTTATTTCTATATGGTGTAGTACAGGAAAGGTTTGTCTATGTTGTAGCAAAATCTGAACGAGCGCTTGGGAT  
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GNTATCTATCTAATCTATCTATATATATACACACAACTCTTATATTTAATTGGTCCCTGAAGTCGCATACATCAATCACAT  
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Q9925893.Seq

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Q9925894.seq

Page 1







Q9925898.Seq

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Q9925899.Seq

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09925900.Seq

Page 1

Q9925901.Seq

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Q9925902.Seq

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Q9925903.seq

Q9925903.seq

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 GCCCAAAANTCCAAGNCATTTGTTATGGATCTAGNTAGAAAAACAGGTGAAAAAGNGTGACCGATTGNGTGNNTTNNTTTACTATAA  
 CGAGTNTTGTCTNGNNGAACCAAGGCTCTGGCATTTGGGGACNTTCAATCCNAAGAAATGNTTGGCCGGGG

T02220 2460T660

MAGPIE - PROJECT - xenopus

GROUP: S10-1

STATE: protein\_dna

-- S10-2 &gt;

Mon Jul 10 08:50:36 EDT 2000

sort by signatureSTATUS | CONTIG INFO | ANNOTATIONS | PRIMERS | ANNOTATOR

The S10-1 group contains 96 contigs.

ID	Bases	N	AA	X	Evidence	Forms	EC	Description	O Rep
S10-1-A1	726 bp	8	241 aa	6	1p 1d 2m	V F M S		> heterogeneous nuclear ribonucleoprotein M (HNRNP M) > dupl	
S10-1-A10	658 bp		219 aa	1	1d 2m	V F M S		> Ribosomal RNA intergenic spacer region	
S10-1-A11	656 bp	1	218 aa	1	1p 1d 2m	V F M S		> citrin/Aralar2 (partial) [CDS CATEGORY]	
S10-1-A12	592 bp	26	197 aa	16	1d 2m	V F M S		> Xenopus EST	
S10-1-A2	698 bp	11	232 aa	12	1d 2m	V F M S		> DNA topoisomerase I (partial) [Helicases]	
S10-1-A3	682 bp	2	227 aa	2	1p 1d 2m	V F M S		> EGF receptor substrate eps15r (FL) [CDS CATEGORY]	
S10-1-A4	765 bp	4	254 aa	4	2p 1d 2m	V F M S		> RNA binding protein (partial) [CDS CATEGORY]	
S10-1-A5	717 bp	1	238 aa	1	1p 1d 2m	V F M S		> tousled-like kinase 1 (FL) [CDS CATEGORY]	
S10-1-A6	451 bp		150 aa	1	2p 1d 2m	V F M S		> histone H3a (partial) [CDS CATEGORY]	
S10-1-A7	718 bp	7	239 aa	7	3p 1d 2m	V F M S		> vector > duplicate [CDS CATEGORY]	
S10-1-A8	718 bp	5	239 aa	5	1d 2m	V F M S		> Xenopus EST	
S10-1-A9	720 bp	4	239 aa	3	1p 1d 2m	V F M S		> Methionine-trna synthetase (partial) [CDS CATEGORY]	
S10-1-B1	720 bp	4	239 aa	3	1p 1d 2m	V F M S		> udp-n-acetylglucosamine pyrophosphorylase (partial) [CDS	
S10-1-B10	650 bp	1	216 aa	2	1p 1d 2m	V F M S		> U2 snrnp-specific A' (FL) [CDS CATEGORY]	

# 7/20/00 9:23 PM

S10-1-B11	660 bp	1	219 aa	1	1p 1d 2m	V F M S
S10-1-B12	639 bp	2	212 aa	2	1p 1d 2m	V F M S
S10-1-B2	728 bp	1	242 aa	2	1p 1d 2m	V F M S
S10-1-B3	763 bp	5	254 aa	3	1p 1d 2m	V F M S
S10-1-B4	710 bp	2	236 aa	3	1p 1d 2m	V F M S
S10-1-B5	687 bp	1	228 aa	1	1p 1d 2m	V F M S
S10-1-B6	721 bp	51	240 aa	42	3p 1d 2m	V F M S
S10-1-B7	704 bp	2	234 aa	3	1d 2m	V F M S
S10-1-B8	709 bp	11	236 aa	10	1d 2m	V F M S
S10-1-B9	644 bp	2	214 aa	2	1p 1d 2m	V F M S
S10-1-C1	695 bp	1	231 aa	2	1p 1d 2m	V F M S
S10-1-C10	724 bp	12	241 aa	11	1p 1d 2m	V F M S
S10-1-C11	667 bp	3	222 aa	3	1d 2m	V F M S
S10-1-C12	668 bp	1	222 aa	2	1p 1d 2m	V F M S
S10-1-C2	713 bp	7	237 aa	7	1p 1d 2m	V F M S
S10-1-C3	716 bp	2	238 aa	2	3p 1d 2m	V F M S
S10-1-C4	675 bp	5	224 aa	4	1d 2m	V F M S
S10-1-C5	716 bp	2	238 aa	2	1p 1d 2m	V F M S
S10-1-C6	717 bp	5	238 aa	4	1p 1d 2m	V F M S
S10-1-C7	709 bp		236 aa	1	1d 2m	V F M S
S10-1-C8	697 bp	2	232 aa	3	2p 1d 2m	V F M S
S10-1-C9	667 bp	1	222 aa	2	1p 1d 2m	V F M S
S10-1-D1	770 bp	6	256 aa	6	1p 1d 2m	V F M S
S10-1-D10	666 bp	1	221 aa	1	1d 2m	V F M S
S10-1-D11	666 bp	2	221 aa	1	3p 1d 2m	V F M S
S10-1-D12	471 bp		156 aa		1d 2m	V F M S
S10-1-D2	695 bp	2	231 aa	2	1p 1d 2m	V F M S
S10-1-D3	714 bp	8	237 aa	7	2p 1d 2m	V F M S
S10-1-D4	716 bp	1	238 aa	2	1p 1d 2m	V F M S
S10-1-D5	718 bp	3	239 aa	2	2p 1d 2m	V F M S

> MAP kinase (FL) [CDS CATEGORY]

> apoptosis inhibitor 5 (FL) [CDS CATEGORY]

> zinc finger protein (partial) [CDS CATEGORY]

> ferritin heavy chain (FL) > Duplicate [CDS CATEGORY]

> mitosis protein DIM1 (FL) [CDS CATEGORY]

> proteasome 26S subunit (partial) [CDS CATEGORY]

> xenopus EST > Duplicate [CDS CATEGORY]

> Xenopus EST [CDS CATEGORY]

> Xenopus EST [CDS CATEGORY]

> k1aa0779 protein (partial) [CDS CATEGORY]

> formaldehyde dehydrogenase (FDH) (partial) [CDS CATEGORY]

> polyubiquitin (FL) > duplicate [CDS CATEGORY]

> Xenopus EST [CDS CATEGORY]

> Rna polymerase II EF ELL2 (FL) > duplicate [RNA polymeras

> DHM1 (FL) [RNA modification]

> Xenopus EST [CDS CATEGORY]

> Xenopus EST

> WD containing protein, conserved (FL) [CDS CATEGORY]

> VON hippel-lindau binding protein 1 (partial) > Duplicate

> Xenopus EST

> gdf-9 like (FL) [CDS CATEGORY]

> ADP/ATP translocase (FL) [CDS CATEGORY]

> NG,NG-dimethylarginine dimethylaminohydrolase (FL) [CDS C

> Xenopus EST

> Xenopus EST [CDS CATEGORY]

> Xenopus EST

> progesterin induced protein (partial) [CDS CATEGORY]

> immediate early response gene 5 like (FL) [CDS CATEGORY]

> cathepsin L (cysteine proteinases) (FL) [CDS CATEGORY]

> melanoma antigen, family D, 1 (FL) [CDS CATEGORY]

# TABLE 4

MAGPIE PROJECT: xenopus GROUP: S10-1

S10-1-D6	717 bp	3	238 aa	2	1p 1d 2m	V F M S
S10-1-D7	707 bp	2	235 aa	2	1d 2m	V F M S
S10-1-D8	710 bp	3	236 aa	3	1d 2m	V F M S
S10-1-D9	665 bp	2	221 aa	2	1d 2m	V F M S
S10-1-E1	721 bp	2	240 aa	3	1d 2m	V F M S
S10-1-E10	720 bp	107	239 aa	65	1p 1d 2m	V F M S
S10-1-E11	664 bp	1	221 aa	2	1d 2m	V F M S
S10-1-E12	661 bp	1	220 aa	2	3p 1d 2m	V F M S
S10-1-E2	571 bp		190 aa	1	1d 2m	V F M S
S10-1-E3	717 bp	2	238 aa	2	1p 1d 2m	V F M S
S10-1-E4	734 bp	168	244 aa	117	1d 2m	V F M S
S10-1-E5	718 bp	3	239 aa	3	1d 2m	V F M S
S10-1-E6	716 bp	1	238 aa	2	2p 1d 2m	V F M S
S10-1-E7	710 bp	5	236 aa	6	1d 2m	V F M S
S10-1-E8	709 bp	1	236 aa	2	1d 2m	V F M S
S10-1-E9	716 bp	4	238 aa	5	3p 1d 2m	V F M S
S10-1-F1	720 bp	3	239 aa	2	3p 1d 2m	V F M S
S10-1-F10	672 bp		223 aa		1p 1d 2m	V F M S
S10-1-F11	638 bp	5	212 aa	5	3p 1d 2m	V F M S
S10-1-F12	641 bp	2	213 aa	2	1p 1d 2m	V F M S
S10-1-F2	723 bp	1	240 aa	1	1d 2m	V F M S
S10-1-F3	715 bp	7	238 aa	7	1p 1d 2m	V F M S
S10-1-F4	707 bp	3	235 aa	4	1d 2m	V F M S
S10-1-F5	716 bp	1	238 aa	2	1p 1d 2m	V F M S
S10-1-F6	801 bp	192	266 aa	139	1d 2m	V F M S
S10-1-F7	705 bp	3	234 aa	2	1d 2m	V F M S
S10-1-F8	705 bp	4	234 aa	4	1p 1d	V F M S
S10-1-F9	653 bp	2	217 aa	2	3p 1d 2m	V F M S
S10-1-G1	720 bp	1	239 aa	1	2p 1d 2m	V F M S

> ATP synthase BETA CHAIN (partial) > Duplicate [CDS CATEGOR
> Xenopus EST [CDS CATEGORY]
> xenopus EST > Duplicate [CDS CATEGORY]
> xenopus EST > Duplicate [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> mRNA capping enzyme 1B (FL) > Duplicate [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> xIRF-6 (partial) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> small nuclear ribonucleoprotein B (partial) > Duplicate [
> Xenopus EST [CDS CATEGORY]
> Xenopus EST
> Xenopus EST [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> cysteine dioxygenase (FL) [CDS CATEGORY]
> emopamil-phenylalkylamine BP FL [CDS CATEGORY]
> RNA Cyclase FL [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> Hypothetical FL [CDS CATEGORY]
> Xenopus EST
> makorin 1 [CDS CATEGORY]
> Xenopus EST
> Xenopus EST [CDS CATEGORY]
> MAP kinase kinase MAPKK (partial) [CDS CATEGORY]
> Selenoprotein like, FL [CDS CATEGORY]
> Aspartate Aminotransferase [CDS CATEGORY]

# TABLE 660

AGPIE PROJECT: xenopus GROUP: S10-1

S10-1-G10	672 bp	10	223 aa	10	1p 1d 2m	V F M S	> membrane-associated diazepam binding inhibitor, MA-DBI FL
S10-1-G11	662 bp		220 aa	1	1p 1d 2m	V F M S	> polyubiquitin (partial) > Duplicate [CDS CATEGORY]
S10-1-G12	718 bp	1	239 aa	1	1p 1d 2m	V F M S	> leucine-rich acidic nuclear phosphoprotein (FL) > Duplicat
S10-1-G2	721 bp	5	240 aa	4	1p 1d 2m	V F M S	> beta-catenin (partial) [CDS CATEGORY]
S10-1-G3	518 bp	3	172 aa	3	3p 1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-1-G4	762 bp	5	253 aa	4	2p 1d 2m	V F M S	> MO25 FL [CDS CATEGORY]
S10-1-G5	697 bp		232 aa	1	2p 1d 2m	V F M S	> ornithine decarboxylase (ODC) (FL) > Duplicate [CDS CATEG
S10-1-G6	713 bp	3	237 aa	4	1d 2m	V F M S	> XFG 5-1/ XFG 5-2 > Duplicate [CDS CATEGORY]
S10-1-G7	707 bp	3	235 aa	4	2p 1d 2m	V F M S	> Protein phosphatase 2 FL [Protein modification]
S10-1-G8	698 bp	9	232 aa	7	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-1-G9	677 bp	2	225 aa	2	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-1-H1	720 bp	5	239 aa	4	3p 1d 2m	V F M S	> Putative kinase (partial) [CDS CATEGORY]
S10-1-H10	658 bp	3	219 aa	3	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-1-H11	336 bp	1	111 aa	1	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-1-H12	675 bp		224 aa		1p 1d 2m	V F M S	> Ran binding protein 11 (partial) [CDS CATEGORY]
S10-1-H2	722 bp	28	240 aa	22	1p 1d 2m	V F M S	> BCL2/adenovirus E1B 19kd-interacting protein 3 [CDS CATEG
S10-1-H3	714 bp	4	237 aa	3	1p 1d 2m	V F M S	> Zyxin Lim Domain protein (partial) [CDS CATEGORY]
S10-1-H4	769 bp	3	256 aa	3	1p 1d 2m	V F M S	> phosphogluconate dehydrogenase (partial) [CDS CATEGORY]
S10-1-H5	776 bp	9	258 aa	7	3p 1d 2m	V F M S	> Possible hydrolase (partial) [CDS CATEGORY]
S10-1-H6	355 bp		118 aa	1	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-1-H7	764 bp	23	254 aa	17	1p 1d 2m	V F M S	> heterogeneous nuclear ribonucleoproteins A2/B1 (FL) > Dupl
S10-1-H8	718 bp	4	239 aa	5	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-1-H9	722 bp	2	240 aa	3	1p 1d 2m	V F M S	> secreted xWnt8 inhibitor sizzled (FL) [CDS CATEGORY]

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Questions and comments about *xenopus* are welcome! Send to [asczyrba@genomes.rockefeller.edu](mailto:asczyrba@genomes.rockefeller.edu)

Questions and comments on *MAGPIE* are welcome! Send to [gaasterland@rockefeller.edu](mailto:gaasterland@rockefeller.edu) [Terry Gaasterland] or [sensencw@niji.imb.nrc.ca](mailto:sensencw@niji.imb.nrc.ca)

TOEZO" E460T650

# M A G P I E - P R O J E C T - xenopus

GROUP: S10-2

STATE: protein dna

&lt;S10-1 -- S10-3 &gt;

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*sort by signature*STATUS | CONTIG INFO | ANNOTATIONS | PRIMERS | ANNOTATOR

The S10-2 group contains 96 contigs.

ID	Bases	N	AA	X	Evidence	Forms	EC	Description	O Re
S10-2-A1	655 bp	1	218 aa	1	1p 1d 2m	V F M S		> glycerol-3-phosphate dehydrogenase > Duplicate [CDS CATEG	
S10-2-A10	686 bp	3	228 aa	4	1d 2m	V F M S		> xenopus EST	
S10-2-A11	681 bp	5	226 aa	4	2p 1d 2m	V F M S		> C elegans ZK546.13 gene product [CDS CATEGORY]	
S10-2-A12	681 bp	4	226 aa	1	3p 1d 2m	V F M S		> sialoglycoprotein (partial) [CDS CATEGORY]	
S10-2-A2	294 bp	33	97 aa	16	1d 2m	V F M S		> Xenopus EST	
S10-2-A3	655 bp		218 aa	1	1p 1d 2m	V F M S		> SWI/SNF related (partial) [CDS CATEGORY]	
S10-2-A4	723 bp	7	240 aa	7	2p 1d 2m	V F M S		> Predicted FL [CDS CATEGORY]	
S10-2-A5	699 bp	2	232 aa	2	1p 1d 2m	V F M S		> geminin H (FL) [CDS CATEGORY]	
S10-2-A6	661 bp	3	220 aa	3	1d 2m	V F M S		> Xenopus EST [CDS CATEGORY]	
S10-2-A7	693 bp	1	230 aa	1	3p 1d 2m	V F M S		> Xenopus EST [CDS CATEGORY]	
S10-2-A8	753 bp	22	250 aa	12	1p 1d 2m	V F M S		> Y BOX binding protein-1 (FL) [CDS CATEGORY]	
S10-2-A9	689 bp	5	229 aa	5	2p 1d 2m	V F M S		> myotubularin 1 (FL) [CDS CATEGORY]	
S10-2-B1	659 bp	1	219 aa	2	3p 1d 2m	V F M S		> Xenopus EST [CDS CATEGORY]	
S10-2-B10	691 bp	1	230 aa	2	2p 1d 2m	V F M S		> pituitary tumor-transforming 1 (FL) [CDS CATEGORY]	



# TABLE 2.4. ESTs

S10-2-B11	667 bp	3	222 aa	3	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-2-B12	798 bp	168	265 aa	129	1d 2m	V F M S	> Xenopus EST
S10-2-B2	625 bp	4	208 aa	5	1p 1d 2m	V F M S	> Putative helicase (partial) [CDS CATEGORY]
S10-2-B3	652 bp		217 aa	1	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-2-B4	655 bp	1	218 aa	1	3p 1d 2m	V F M S	> Putative nexin [CDS CATEGORY]
S10-2-B5	391 bp	1	130 aa	2	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-2-B6	697 bp		232 aa	1	1p 1d 2m	V F M S	> Fatvg (partial) [CDS CATEGORY]
S10-2-B7	692 bp	2	230 aa	2	1p 1d 2m	V F M S	> glycosyl transferase (partial) [CDS CATEGORY]
S10-2-B8	668 bp	4	222 aa	4	1p 1d 2m	V F M S	> RING finger protein (partial) [CDS CATEGORY]
S10-2-B9	721 bp	9	240 aa	9	3p 1d 2m	V F M S	> vector > Duplicate [CDS CATEGORY]
S10-2-C1	673 bp	5	224 aa	6	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-2-C10	692 bp	2	230 aa	2	1d 2m	V F M S	> Xenopus EST
S10-2-C11	683 bp	2	227 aa	3	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-2-C12	680 bp		226 aa	1	1d	V F M S	> Xenopus EST
S10-2-C2	670 bp	3	223 aa	3	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-2-C3	670 bp	4	223 aa	5	3p 1d 2m	V F M S	> Predicted C elegans homolog, rat exo84 (FL) [CDS CATEGORY]
S10-2-C4	669 bp	8	222 aa	6	1p 1d 2m	V F M S	> eukaryotic translation initiation factor 2G (partial) [CD
S10-2-C5	771 bp	32	256 aa	19	1d 2m	V F M S	> Xenopus EST
S10-2-C6	692 bp	34	230 aa	31	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-2-C7	757 bp	13	252 aa	6	2p 1d 2m	V F M S	> Putative Arabidopsis homologue [CDS CATEGORY]
S10-2-C8	689 bp	4	229 aa	3	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-2-C9	692 bp	2	230 aa	1	1p 1d 2m	V F M S	> ubiquitin-like protein smt3a (FL) > duplicate [CDS Catego
S10-2-D1	671 bp	1	223 aa	2	3p 1d 2m	V F M S	> Putative spicing factor [CDS CATEGORY]
S10-2-D10	691 bp		230 aa	1	1p 1d 2m	V F M S	> ubiquitin-like protein smt3a (FL) > Duplicate [CDS Catego
S10-2-D11	630 bp	5	209 aa	4	2p 1d 2m	V F M S	> lamin B1 (FL) [CDS CATEGORY]
S10-2-D12	668 bp	4	222 aa	5	3p 1d 2m	V F M S	> Xenopus EST > Duplicate [CDS CATEGORY]
S10-2-D2	670 bp	3	223 aa	3	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-2-D3	652 bp	5	217 aa	6	1p 1d 2m	V F M S	> small nuclear ribonucleoprotein polypeptide B" [CDS CATE
S10-2-D4	652 bp	2	217 aa	3	2p 1d 2m	V F M S	> ODC (FL) > Duplicate [CDS CATEGORY]
S10-2-D5	700 bp	2	233 aa	3	1p 1d 2m	V F M S	> snap-25 interacting protein hrs-2 (FL) [CDS CATEGORY]



## TABLE 240

S10-2-D6	688 bp		229 aa	1	1p 1d 2m	V F M S
S10-2-D7	657 bp	1	218 aa	1	1d 2m	V F M S
S10-2-D8	687 bp	6	228 aa	6	1d 2m	V F M S
S10-2-D9	691 bp		230 aa	1	1p 1d 2m	V F M S
S10-2-E1	620 bp	1	206 aa	2	1d 2m	V F M S
S10-2-E10	686 bp	2	228 aa	1	1d 2m	V F M S
S10-2-E11	679 bp	1	226 aa	2	1d 2m	V F M S
S10-2-E12	768 bp	24	255 aa	14	2p 1d 2m	V F M S
S10-2-E2	640 bp	2	213 aa	2	1d 2m	V F M S
S10-2-E3	469 bp	51	156 aa	34	2p 1d 2m	V F M S
S10-2-E4	722 bp		240 aa	1	1p 1d 2m	V F M S
S10-2-E5	698 bp		232 aa	1	1d 2m	V F M S
S10-2-E6	700 bp	2	233 aa	3	3p 1d 2m	V F M S
S10-2-E7	687 bp	3	228 aa	1	1p 1d 2m	V F M S
S10-2-E8	688 bp	10	229 aa	8	1p 1d 2m	V F M S
S10-2-E9	690 bp	1	229 aa	1	2p 1d 2m	V F M S
S10-2-F1	662 bp		220 aa	1	3p 1d 2m	V F M S
S10-2-F10	674 bp	1	224 aa	2	3p 1d 2m	V F M S
S10-2-F11	639 bp	3	212 aa	1	1p 1d 2m	V F M S
S10-2-F12	675 bp	3	224 aa	3	1d 2m	V F M S
S10-2-F2	648 bp		215 aa		1p 1d 2m	V F M S
S10-2-F3	661 bp	56	220 aa	46	1d 2m	V F M S
S10-2-F4	661 bp	1	220 aa	2	1p 1d 2m	V F M S
S10-2-F5	748 bp	1	249 aa	2	1p 1d 2m	V F M S
S10-2-F6	752 bp	5	250 aa	6	1p 1d 2m	V F M S
S10-2-F7	690 bp	4	229 aa	3	1p 1d 2m	V F M S
S10-2-F8	661 bp	2	220 aa	3	1d 2m	V F M S
S10-2-F9	687 bp	2	228 aa	1	2p 1d 2m	V F M S
S10-2-G1	649 bp	12	216 aa	8	1p 1d 2m	V F M S

> Conserved protein KIAA0007 (partial) [CDS CATEGORY]
> Xenopus EST
> Xenopus EST [CDS CATEGORY]
> Predicted conserved protein (partial) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> conserved nucleolar protein (KKE/D repeat) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> cytochrome c oxidase subunit I (nFL) > duplicate [CDS CAT
> glycerol-3-phosphate dehydrogenase (FL) > Duplicate [CDS
> Xenopus EST [CDS CATEGORY]
> Xenopus EST > myosin heavy chain homology [CDS CATEGORY]
> translocon-associated Protein, BETA subunit precursor (FL)
> RNA binding protein hnRNP- D/nrp-1b [CDS CATEGORY]
> Conserved protein homology (FL) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> Predicted conserved protein (FL) [CDS CATEGORY]
> NADH dehydrogenase subunit 2 (partial) [CDS CATEGORY]
> Xenopus EST
> HEAT shock cognate 71 KD protein (partial) [CDS CATEGORY]
> Xenopus EST
> glutamine synthetase (nFL) [CDS CATEGORY]
> leucine-rich acidic nuclear protein (FL) > Duplicate [CDS
> transcription factor (TFIIIC) alpha chain (partial) [CDS
> granulin precursor (nFL) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> MutT similarity [CDS CATEGORY]
> mRNA capping enzyme (FL) > Duplicate [CDS CATEGORY]

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S10-2-G10	687 bp	1	228 aa	1	1p 1d 2m	V F M S	> CREATINE KINASE, B CHAIN [CDS CATEGORY]
S10-2-G11	826 bp	202	275 aa	141	1d 2m	V F M S	> Vector
S10-2-G12	659 bp	5	219 aa	5	2p 1d 2m	V F M S	> Putative Sodium/phosphate transporter (nFL) > Xenopus EST
S10-2-G2	676 bp	1	225 aa	1	1p 1d 2m	V F M S	> ferritin heavy chain 1 (FL) [CDS CATEGORY]
S10-2-G3	676 bp		225 aa	1	1p 1d 2m	V F M S	> kina0780 protein (partial) [CDS CATEGORY]
S10-2-G4	664 bp	2	221 aa	2	3p 1d 2m	V F M S	> TFIIIS elongation factor (FL) > Duplicate [CDS CATEGORY]
S10-2-G5	720 bp	5	239 aa	3	1p 1d 2m	V F M S	> All-1 related protein (partial) [CDS CATEGORY]
S10-2-G6	692 bp	6	230 aa	6	1d 2m	V F M S	> Xenopus EST
S10-2-G7	686 bp	3	228 aa	4	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-2-G8	664 bp		221 aa	1	1d 2m	V F M S	> Xenopus EST
S10-2-G9	690 bp	3	229 aa	3	3p 1d 2m	V F M S	> succinyl-CoA synthetase beta subunit (FL) [CDS CATEGORY]
S10-2-H1	677 bp	1	225 aa	2	1d 2m	V F M S	> XFG 5-1 - XFG 5-2 zinc finger proteins (partial) > Duplica
S10-2-H10	687 bp	4	228 aa	3	1p 1d 2m	V F M S	> Isocitrate dehydrogenase subunit ALPHA (nFL) [CDS CATEGORY]
S10-2-H11	658 bp	2	219 aa	2	3p 1d 2m	V F M S	> protein phosphatase 2, regulatory subunit B (FL) [CDS CAT
S10-2-H12	680 bp	1	226 aa	2	3p 1d 2m	V F M S	> topoisomerase-related function protein (FL) [CDS CATEGORY]
S10-2-H2	724 bp	5	241 aa	4	1p 1d 2m	V F M S	> g-alpha-q protein - african clawed frog (partial) [CDS CA
S10-2-H3	675 bp	1	224 aa	1	2p 1d 2m	V F M S	> STE20-like kinase 3 (FL) [CDS CATEGORY]
S10-2-H4	725 bp	5	241 aa	5		V F M S	> MAP kinase activated (?) > Duplicate [CDS CATEGORY]
S10-2-H5	696 bp	3	231 aa	3	1p 1d 2m	V F M S	> pyruvate dehydrogenase kinase (FL) [CDS CATEGORY]
S10-2-H6	659 bp	5	219 aa	5	2p 1d 2m	V F M S	> secretory carrier membrane protein 1 (nFL) [CDS CATEGORY]
S10-2-H7	687 bp	4	228 aa	4	1p 1d 2m	V F M S	> rna-binding protein (KH domain) (FL) [CDS CATEGORY]
S10-2-H8	342 bp	1	113 aa	1	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-2-H9	689 bp	5	229 aa	6	1p 1d 2m	V F M S	> leucine-rich acidic nuclear protein (FL) > Duplicate [CDS

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Questions and comments about *xenopus* are welcome! Send to [asczyrba@genomes.rockefeller.edu](mailto:asczyrba@genomes.rockefeller.edu)

Questions and comments on *MAGPIE* are welcome! Send to [gaasterland@rockefeller.edu](mailto:gaasterland@rockefeller.edu) [Terry Gaasterland] or [sensencw@niji.imb.nrc.ca](mailto:sensencw@niji.imb.nrc.ca)

# MAGPIE - PROJECT - xenopus

GROUP: S10-3

STATE: protein\_dna

<S10-2 -- S10-4 >

Mon Jul 10 08:50:36 EDT 2000

sort by signature

STATUS | CONTIG INFO | ANNOTATIONS | PRIMERS | ANNOTATOR

The S10-3 group contains 96 contigs.

ID	Bases	N	AA	X	Evidence	Forms	EC	Description	OR
S10-3-A1	662 bp	3	220 aa	4	1d 2m	V F M S		> Xenopus EST	
S10-3-A10	639 bp	6	212 aa	5	3p 1d 2m	V F M S		> 18S ribosomal RNA [CDS CATEGORY]	
S10-3-A11	639 bp	5	212 aa	4	1p 1d 2m	V F M S		> von hippel-lindau binding protein 1 (FL) > xenopus EST > D	
S10-3-A12	636 bp	3	211 aa	2	1p 1d 2m	V F M S		> small nuclear ribonucleoprotein B' (FL) > Duplicate [CDS	
S10-3-A2	664 bp	3	221 aa	3	3p 1d 2m	V F M S		> nuclear Y/CCAAT-box binding factor A subunit NF-YA (FL)	
S10-3-A3	642 bp		213 aa		3p 1d 2m	V F M S		> co-factor nherf protein (partial) > xenopus EST [CDS CATE	
S10-3-A4	658 bp		219 aa	1	1p 1d 2m	V F M S		> kiaa0515 protein (partial) > xenopus EST [CDS CATEGORY]	
S10-3-A5	655 bp		218 aa	1	1p 1d 2m	V F M S		> kiaa1109 protein (partial) [CDS CATEGORY]	
S10-3-A6	655 bp	1	218 aa	1	1p 1d 2m	V F M S		> heterogeneous nuclear ribonucleoprotein C (FL) [CDS CATEG	
S10-3-A7	653 bp		217 aa	1	3p 1d 2m	V F M S		> Elongation factor 3 (nFL) [CDS CATEGORY]	
S10-3-A8	656 bp		218 aa	1	1p 1d 2m	V F M S		> G2/MITOTIC-SPECIFIC cyclin B1 (FL) > Duplicate [CDS CATEG	
S10-3-A9	656 bp	9	218 aa	6	2p 1d 2m	V F M S		> Sphere organelles protein 1 (FL) [CDS CATEGORY]	
S10-3-B1	470 bp	2	156 aa	3	1d 2m	V F M S		> Xenopus EST [CDS CATEGORY]	
S10-3-B10	651 bp	3	216 aa	3	3p 1d 2m	V F M S		> transmembrane glycoprotein (nFL) > xenopus EST [CDS CATEG	

# TABLE 20-2460T550

S10-3-B11	599 bp	199 aa	1	1p 1d 2m	V F M S	> putative GTP-ase activating protein for Arf > Xenopus EST
S10-3-B12	636 bp	211 aa		3p 1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-3-B2	630 bp	209 aa	2	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-3-B3	658 bp	219 aa	2	1d 2m	V F M S	> XFG 5-1 and XFG 5-2 > Xenopus EST [CDS CATEGORY]
S10-3-B4	662 bp	220 aa	2	1p 1d 2m	V F M S	> Putative transcription factor (partial) [CDS CATEGORY]
S10-3-B5	658 bp	219 aa	1	1p 1d 2m	V F M S	> nuclear RNA helicase (nFL) [CDS CATEGORY]
S10-3-B6	475 bp	158 aa	6	1d 2m	V F M S	> alpha e-catenin (partial) [CDS CATEGORY]
S10-3-B7	702 bp	233 aa	1	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-3-B8	653 bp	217 aa	1	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-3-B9	649 bp	216 aa	3	3p 1d 2m	V F M S	> xenopus EST > Duplicate [CDS CATEGORY]
S10-3-C1	713 bp	237 aa	3	1p 1d 2m	V F M S	> ser-thr protein phosphatase (FL) > Duplicate [CDS CATEGOR
S10-3-C10	627 bp	208 aa		3p 1d 2m	V F M S	> cdc47-2p (FL) > Duplicate [CDS CATEGORY]
S10-3-C11	634 bp	211 aa	2	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-3-C12	636 bp	211 aa	1	1p 1d 2m	V F M S	> Hypothetical protein (FL) [CDS CATEGORY]
S10-3-C2	612 bp	203 aa	10	1d 2m	V F M S	> Xenopus EST
S10-3-C3	662 bp	220 aa	3	1p 1d	V F M S	> PITSLRE protein kinases p34cdc2 (FL) [CDS CATEGORY]
S10-3-C4	664 bp	221 aa	2	1p 1d 2m	V F M S	> TPR-containing protein (partial) [CDS CATEGORY]
S10-3-C5	657 bp	218 aa	4	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-3-C6	659 bp	219 aa	4	1p 1d 2m	V F M S	> G2/MITOTIC-SPECIFIC cyclin B2 (FL) > Duplicate [CDS CATEG
S10-3-C7	659 bp	219 aa	1	1p 1d 2m	V F M S	> PA26-T2 nuclear protein (nFL) [CDS CATEGORY]
S10-3-C8	638 bp	212 aa	2	1p 1d 2m	V F M S	> KIAA0064 (FL) [CDS CATEGORY]
S10-3-C9	630 bp	209 aa	2	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-3-D1	31 bp	10 aa	1		V F M S	> No insert
S10-3-D10	630 bp	209 aa	29	1p 1d 2m	V F M S	> kiaa0544 protein (nFL) [CDS CATEGORY]
S10-3-D11	665 bp	221 aa	62	1d 2m	V F M S	> Xenopus EST
S10-3-D12	613 bp	204 aa	1	1p 1d 2m	V F M S	> Focal adhesion kinase 1 (FADK 1) (partial) [CDS CATEGORY]
S10-3-D2	665 bp	221 aa	3	3p 1d 2m	V F M S	> eukaryotic translation initiation factor 4B (partial) > Xe
S10-3-D3	662 bp	220 aa	4	1p 1d 2m	V F M S	> osteoactivin (partial) > xenopus EST [CDS CATEGORY]
S10-3-D4	664 bp	221 aa	4	1p 1d 2m	V F M S	> guanine nucleotide-binding protein G(I) (partial) [CDS CA
S10-3-D5	661 bp	220 aa	1	1p 1d 2m	V F M S	> Xbtg1 (FL) [CDS CATEGORY]

# TABLE 4: ESTs

S10-3-D6	658 bp		219 aa	1	1p 1d 2m	V F M S
S10-3-D7	650 bp	1	216 aa	2	1d 2m	V F M S
S10-3-D8	655 bp	2	218 aa	2	1p 1d 2m	V F M S
S10-3-D9	647 bp	2	215 aa	3	1p 1d 2m	V F M S
S10-3-E1	720 bp	11	239 aa	9	3p 1d 2m	V F M S
S10-3-E10	609 bp	2	202 aa	2	1p 1d 2m	V F M S
S10-3-E11	750 bp	104	249 aa	74	1d 2m	V F M S
S10-3-E12	751 bp	127	250 aa	93	1d 2m	V F M S
S10-3-E2	664 bp	2	221 aa	2	1p 1d 2m	V F M S
S10-3-E3	665 bp	10	221 aa	10	1d 2m	V F M S
S10-3-E4	658 bp	1	219 aa	2	1p 1d 2m	V F M S
S10-3-E5	657 bp	1	218 aa	1	1d 2m	V F M S
S10-3-E6	663 bp	1	220 aa	1	1p 1d 2m	V F M S
S10-3-E7	640 bp	1	213 aa	2	1p 1d 2m	V F M S
S10-3-E8	650 bp	1	216 aa	2	2p 1d 2m	V F M S
S10-3-E9	361 bp		120 aa	1	1d 2m	V F M S
S10-3-F1	667 bp	2	222 aa	3	1p 1d 2m	V F M S
S10-3-F10	625 bp	3	208 aa	3	1p 1d 2m	V F M S
S10-3-F11	611 bp	1	203 aa	1	1p 1d 2m	V F M S
S10-3-F12	636 bp	1	211 aa	1	1p 1d 2m	V F M S
S10-3-F2	632 bp	3	210 aa	4	2p 1d 2m	V F M S
S10-3-F3	665 bp	2	221 aa	2	1p 1d 2m	V F M S
S10-3-F4	664 bp	3	221 aa	3	2p 1d 2m	V F M S
S10-3-F5	651 bp	1	216 aa	1	1p 1d 2m	V F M S
S10-3-F6	659 bp		219 aa	1	1p 1d 2m	V F M S
S10-3-F7	658 bp		219 aa	1	1d 2m	V F M S
S10-3-F8	658 bp	2	219 aa	2	1p 1d 2m	V F M S
S10-3-F9	558 bp		185 aa		1d 2m	V F M S
S10-3-G1	627 bp	5	208 aa	5	1p 1d 2m	V F M S

> xrad51 (FL) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> protein phosphatase 2C beta (FL) > Duplicate [CDS CATEGOR
> G2/MITOTIC-SPECIFIC cyclin B2 (FL) > Duplicate [CDS CATEG
> vector > Duplicate [CDS CATEGORY]
> XICdc1 (FL) [CDS CATEGORY]
> Xenopus EST
> Xenopus EST
> homeobox transcription factor iriouis 3 (FL) [CDS Catego
> Xenopus EST [CDS CATEGORY]
> splicing factor, arginine/serine-rich 7 (FL) [CDS CATEGOR
> xenopus EST > Duplicate [CDS CATEGORY]
> xbr-1b/Vox-1 (FL) [CDS CATEGORY]
> cyclin ania-6a (partial) [CDS CATEGORY]
> chromodomain helicase DNA binding protein 3 (partial) [CD
> Xenopus EST [CDS CATEGORY]
> aspartate AMINOTRANSFERASE (FL) [CDS CATEGORY]
> hnmp A1 (FL) > Duplicate [CDS CATEGORY]
> nucleoporin p54 (FL) > Xenopus EST [CDS CATEGORY]
> STE20/KIAA1264 protein kinase > Xenopus EST [CDS CATEGORY
> C elegans [CDS CATEGORY]
> Arg/Ser-rich 6 splicing factor 6 (partial) [CDS CATEGORY]
> histone stem-loop binding protein (FL) [CDS CATEGORY]
> Human orf hspc017 (partial) > Xenopus EST [CDS CATEGORY]
> transcription EF ELL gene (FL) > xenopus EST > Duplicate
> ribosomal protein S6 > Xenopus EST [CDS CATEGORY]
> xIRF-6 (FL) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> CGI-23 protein (partial) [CDS CATEGORY]



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S10-3-G10	603 bp	2	200 aa	2	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-3-G11	643 bp	23	214 aa	20	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-3-G12	639 bp	3	212 aa	3	2p 1d 2m	V F M S	> ribonucleoside-diphosphate reductase (FL) > Xenopus EST [
S10-3-G2	665 bp	4	221 aa	3	3p 1d 2m	V F M S	> Ankyrin like protein [CDS CATEGORY]
S10-3-G3	628 bp	1	209 aa	2	2p 1d 2m	V F M S	> COP9 complex subunit 3 (FL) [CDS CATEGORY]
S10-3-G4	662 bp	2	220 aa	3	1p 1d 2m	V F M S	> fibrillarin (nFL) [CDS CATEGORY]
S10-3-G5	658 bp	1	219 aa	2	1p 1d 2m	V F M S	> DNA topoisomerase II (partial) [CDS CATEGORY]
S10-3-G6	660 bp	2	219 aa	2	2p 1d 2m	V F M S	> hnnp G (FL) > xenopus EST > Duplicate [CDS CATEGORY]
S10-3-G7	654 bp	2	217 aa	2	2p 1d 2m	V F M S	> Ser/Thr phosphatase pp2a-4 (FL) > Duplicate [CDS CATEGORY]
S10-3-G8	654 bp		217 aa		1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-3-G9	645 bp		214 aa		1p 1d 2m	V F M S	> Stromal antigen 2 SA-2 (partial) [CDS CATEGORY]
S10-3-H1	662 bp	18	220 aa	16	1d 2m	V F M S	> xenopus EST > Ribosomal RNA [CDS CATEGORY]
S10-3-H10	641 bp		213 aa	1	1p 1d 2m	V F M S	> Rac/Rho cdc42 (FL) > Duplicate [CDS CATEGORY]
S10-3-H11	641 bp		213 aa	1	2p 1d 2m	V F M S	> G2/Mitotic specific cyclin B (FL) > xenopus EST > Duplicat
S10-3-H12	621 bp		206 aa		1p 1d 2m	V F M S	> p24 delta1 putative cargo receptor (nFL) > Duplicate [CDS
S10-3-H2	652 bp	2	217 aa	3	1d 2m	V F M S	> xenopus EST > Duplicate [CDS CATEGORY]
S10-3-H3	660 bp		219 aa		1p 1d 2m	V F M S	> kiaa0290 cdc15 gas-7 (FL) [CDS CATEGORY]
S10-3-H4	710 bp		236 aa	1	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-3-H5	655 bp	1	218 aa	1	1d 2m	V F M S	> Xenopus EST
S10-3-H6	659 bp	1	219 aa	2	1p 1d 2m	V F M S	> EMK/Mark 2 protein kinase [CDS CATEGORY]
S10-3-H7	657 bp	1	218 aa	1	1p 1d 2m	V F M S	> eps8 binding protein (FL) [CDS CATEGORY]
S10-3-H8	655 bp		218 aa	1	2p 1d 2m	V F M S	> putative progesterone-binding protein (FL) > xenopus EST
S10-3-H9	645 bp		214 aa		1p 1d 2m	V F M S	> MDM-2 c (partial) [CDS CATEGORY]

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Questions and comments on MAGPIE are welcome! Send to [gaasterland@rockefeller.edu](mailto:gaasterland@rockefeller.edu) [Terry Gaasterland] or [sensencw@niji.imb.nrc.ca](mailto:sensencw@niji.imb.nrc.ca)

# MAGPIE - PROJECT - xenopus

GROUP: S10-4

STATE: protein\_dna

<S10-3 -- S10-5 >

Mon Jul 10 08:50:36 EDT 2000

sort by signature

STATUS | CONTIG INFO | ANNOTATIONS | PRIMERS | ANNOTATOR

The S10-4 group contains 96 contigs.

ID	Bases	N	AA	X	Evidence	Forms	EC	Description	Oth Rep
S10-4-A1	616 bp	2	205 aa	2	1d 2m	V F M S		> Xenopus EST [CDS CATEGORY]	
S10-4-A10	690 bp	5	229 aa	5	2p 1d 2m	V F M S		> Human orf (FL) [CDS CATEGORY]	
S10-4-A11	682 bp	3	227 aa	4	1p 1d 2m	V F M S		> purinergic receptor (partial) [CDS CATEGORY]	
S10-4-A12	688 bp		229 aa	1	1p 1d 2m	V F M S		> hmg-1 (partial) > xenopus EST > duplicate [CDS CATEGORY]	
S10-4-A2	636 bp	3	211 aa	3	1p 1d 2m	V F M S		> poly a binding protein (partial) [CDS CATEGORY]	
S10-4-A3	600 bp	2	199 aa	2	1p 1d 2m	V F M S		> 40S ribosomal protein S4 (nFL) > Duplicate [CDS CATEGORY]	
S10-4-A4	633 bp	1	210 aa	1	1p 1d 2m	V F M S		> small nuclear ribonucleoprotein B' (FL) > Duplicate [CDS	
S10-4-A5	701 bp	3	233 aa	3	1p 1d 2m	V F M S		> Translation initiation factor Eif1 (FL) [CDS CATEGORY]	
S10-4-A6	686 bp	2	228 aa	3	1p 1d 2m	V F M S		> holocytochrome c synthetase (nFL) > xenopus EST [CDS CATE	
S10-4-A7	747 bp	6	248 aa	6	1d 2m	V F M S		> Xenopus EST > Xenopus REM sequence [CDS CATEGORY]	
S10-4-A8	646 bp	12	215 aa	10	1d 2m	V F M S		> dsrna-binding protein 4F.1 (FL) > xenopus EST > duplicate	
S10-4-A9	694 bp	4	231 aa	3	1p 1d 2m	V F M S		> xAN11 wd-repeat protein (FL) [CDS CATEGORY]	
S10-4-B1	563 bp	1	187 aa	2	1d 2m	V F M S		> Xenopus EST	
S10-4-B10	693 bp	4	230 aa	3	1p 1d 2m	V F M S		> MRP atpase (nFL) > xenopus EST [CDS CATEGORY]	

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S10-4-B11	689 bp	3	229 aa	4	3p 1d 2m	V F M S	> Xenopus EST > Yeast/elegans hypothetical (FL) [CDS CATEGORY]
S10-4-B12	720 bp	9	239 aa	9	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-4-B2	646 bp	2	215 aa	3	1p 1d 2m	V F M S	> dsrna-binding protein 4F.1 (FL) > xenopus EST > duplicate
S10-4-B3	631 bp		210 aa	1	1p 1d 2m	V F M S	> nm23-x7 (FL) > xenopus EST [CDS CATEGORY]
S10-4-B4	619 bp		206 aa	1	2p 1d 2m	V F M S	> genethonin 1 (FL) > Duplicate (partial) [CDS CATEGORY]
S10-4-B5	703 bp	5	234 aa	5	2p 1d 2m	V F M S	> glucose/oxygen regulated protein (partial) [CDS CATEGORY]
S10-4-B6	702 bp	2	233 aa	1	1d 2m	V F M S	> xenopus EST > Duplicate [CDS CATEGORY]
S10-4-B7	696 bp	1	231 aa	1	2p 1d 2m	V F M S	> Fused toes/ Fif protein (FL) [CDS CATEGORY]
S10-4-B8	693 bp	6	230 aa	6	1p 1d 2m	V F M S	> hsp-90 (FL) > xenopus EST > Duplicate [CDS CATEGORY]
S10-4-B9	695 bp	40	231 aa	17	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-4-C1	645 bp	7	214 aa	6	1d 2m	V F M S	> Xenopus EST
S10-4-C10	334 bp	3	111 aa	4	1d	V F M S	> Xenopus EST
S10-4-C11	689 bp	2	229 aa	2	1p 1d 2m	V F M S	> dolichyl-phosphate beta-glucosyltransferase (partial) [CD
S10-4-C12	688 bp	46	229 aa	42	1p 1d 2m	V F M S	> rna-binding protein xlnrnp1 (FL) [CDS CATEGORY]
S10-4-C2	642 bp		213 aa		1p 1d 2m	V F M S	> xGATA-4 (FL) [CDS CATEGORY]
S10-4-C3	630 bp		209 aa		1p 1d 2m	V F M S	> MCT-3 (partial) [CDS CATEGORY]
S10-4-C4	584 bp		194 aa	1	1p 1d 2m	V F M S	> dmr-n9 protein (partial) [CDS CATEGORY]
S10-4-C5	690 bp	6	229 aa	6	1p 1d 2m	V F M S	> claudin-6 (FL) > duplicate [CDS CATEGORY]
S10-4-C6	704 bp	6	234 aa	5	1d 2m	V F M S	> Xenopus EST
S10-4-C7	680 bp	5	226 aa	4	1p 1d 2m	V F M S	> guanine nucleotide-binding protein G(I), alpha-1 subunit (
S10-4-C8	693 bp	5	230 aa	3	1p 1d 2m	V F M S	> Hyaluronidase (FL) [CDS CATEGORY]
S10-4-C9	691 bp	1	230 aa	2	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-4-D1	643 bp	1	214 aa	2	1d 2m	V F M S	> Xenopus EST
S10-4-D10	696 bp	7	231 aa	6	1d 2m	V F M S	> xenopus EST > Duplicate [CDS CATEGORY]
S10-4-D11	693 bp	14	230 aa	13	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-4-D12	386 bp	1	128 aa	2	1d	V F M S	> Xenopus EST
S10-4-D2	642 bp	1	213 aa	1	1p 1d 2m	V F M S	> protein arginine n-methyltransferase 1 (FL) [CDS CATEGORY]
S10-4-D3	632 bp		210 aa	1	2p 1d 2m	V F M S	> met-10+ protein (nFL) > xenopus EST [CDS CATEGORY]
S10-4-D4	632 bp	1	210 aa	2	1p 1d 2m	V F M S	> gt334 protein (partial) [CDS CATEGORY]
S10-4-D5	686 bp	1	228 aa	2	2p 1d 2m	V F M S	> beta-amyloid precursor protein (FL) [CDS CATEGORY]



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S10-4-D6	703 bp	5	234 aa	6	1p 1d 2m	V F M S
S10-4-D7	665 bp	2	221 aa	2	1d 2m	V F M S
S10-4-D8	695 bp	4	231 aa	3	3p 1d 2m	V F M S
S10-4-D9	692 bp	1	230 aa	1	3p 1d 2m	V F M S
S10-4-E1	644 bp	3	214 aa	3	1p 1d 2m	V F M S
S10-4-E10	696 bp	6	231 aa	5	1p 1d 2m	V F M S
S10-4-E11	654 bp	1	217 aa	1	1d 2m	V F M S
S10-4-E12	699 bp	19	232 aa	17	1p 1d 2m	V F M S
S10-4-E2	642 bp	10	213 aa	10	2p 1d 2m	V F M S
S10-4-E3	631 bp	1	210 aa	2	1d 2m	V F M S
S10-4-E4	633 bp		210 aa		1p 1d 2m	V F M S
S10-4-E5	702 bp	4	233 aa	3	1d 2m	V F M S
S10-4-E6	703 bp	4	234 aa	5	3p 1d 2m	V F M S
S10-4-E7	690 bp	1	229 aa	1	1d 2m	V F M S
S10-4-E8	688 bp	5	229 aa	5	1d 2m	V F M S
S10-4-E9	691 bp		230 aa	1	2p 1d 2m	V F M S
S10-4-F1	649 bp	19	216 aa	16	1d 2m	V F M S
S10-4-F10	714 bp	4	237 aa	4	3p 1d 2m	V F M S
S10-4-F11	689 bp	3	229 aa	3	3p 1d 2m	V F M S
S10-4-F12	457 bp		152 aa	1	1d 2m	V F M S
S10-4-F2	148 bp		49 aa	1	1d 2m	V F M S
S10-4-F3	608 bp		202 aa	1	1d 2m	V F M S
S10-4-F4	439 bp	21	146 aa	13	1p 1d 2m	V F M S
S10-4-F5	689 bp		229 aa	1	3p 1d 2m	V F M S
S10-4-F6	701 bp	3	233 aa	4	2p 1d 2m	V F M S
S10-4-F7	694 bp	2	231 aa	3	1p 1d 2m	V F M S
S10-4-F8	694 bp	1	231 aa	2	1d 2m	V F M S
S10-4-F9	691 bp	2	230 aa	3	1d 2m	V F M S
S10-4-G1	647 bp	31	215 aa	29	3p 1d 2m	V F M S

  

> elongation factor 1 gamma (FL) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> Xenopus EST (partial) [CDS CATEGORY]
> xenopus EST > Duplicate [CDS CATEGORY]
> tetratricopeptide repeat domain 4 (FL) [CDS CATEGORY]
> progesterone membrane binding protein (partial) > xenopus
> Xenopus EST [CDS CATEGORY]
> alpha-2 tubulin (FL) [CDS CATEGORY]
> histone binding N1/N2 (FL) > duplicate [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> xRanbp1 (FL) [CDS CATEGORY]
> Mitochondrial > Similar [CDS CATEGORY]
> ankyrin (brank-2) like (partial) [CDS CATEGORY]
> Xenopus EST
> Xenopus EST [CDS CATEGORY]
> Ca2+-transporting atpase (partial) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> vector > Duplicate [CDS CATEGORY]
> procollagen/acetyltransferase similarity > Duplicate [CDS
> Xenopus EST [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> signal recognition particle SRP-9 (FL) [CDS CATEGORY]
> procollagen/acetyltransferase similarity > Duplicate [CDS
> BAT4 (FL) [CDS CATEGORY]
> hsp-90 (FL) > Duplicate [CDS CATEGORY]
> xenopus repetitive element > xenopus EST > duplicate [CDS
> Xenopus EST
> Xenopus EST > RPB-25 (FL) [CDS CATEGORY]

S10-4-G10	691 bp	3	230 aa	4	1p 1d 2m	V F M S	> calumenin (FL) [CDS CATEGORY]
S10-4-G11	734 bp	47	244 aa	39	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-4-G12	671 bp	35	223 aa	28	1p 1d 2m	V F M S	> tfiid 18K subunit TAF (nFL) > xenopus EST [CDS CATEGORY]
S10-4-G2	625 bp		208 aa	1	1p 1d 2m	V F M S	> RRM RNA binding protein gry-rbp (FL) > Duplicate [CDS CAT
S10-4-G3	639 bp		212 aa		1d 2m	V F M S	> xNopp180 nucleolar phosphoprotein (nFL) [CDS CATEGORY]
S10-4-G4	718 bp	2	239 aa	2	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-4-G5	701 bp	3	233 aa	4	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-4-G6	641 bp	2	213 aa	3	1p 1d 2m	V F M S	> Ca2+/calmodulin-dependent protein kinase II gamma-c (parti
S10-4-G7	696 bp	1	231 aa		1p 1d 2m	V F M S	> peptidylprolyl isomerase D (cyclophilin D) (FL) [CDS CATE
S10-4-G8	677 bp	1	225 aa	2	1p 1d 2m	V F M S	> zinc finger protein (FL) [CDS CATEGORY]
S10-4-G9	691 bp	4	230 aa	4	2p 1d 2m	V F M S	> p24 delta1 putative cargo receptor (partial) > Duplicate
S10-4-H1	708 bp	404	235 aa	149	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-4-H10	673 bp	2	224 aa	2	1p 1d 2m	V F M S	> cold-inducible rna-binding protein (partial) > Duplicate
S10-4-H11	688 bp	4	229 aa	4	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-4-H12	650 bp		216 aa	1	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-4-H2	642 bp	2	213 aa	2	1p 1d 2m	V F M S	> HMG-2 (HMG-2) (partial) [CDS CATEGORY]
S10-4-H3	634 bp		211 aa	1	3p 1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-4-H4	635 bp		211 aa	1	1p 1d 2m	V F M S	> CGI-66 (partial) [CDS CATEGORY]
S10-4-H5	696 bp	4	231 aa	3	3p 1d 2m	V F M S	> CGI-78 protein (FL) [CDS CATEGORY]
S10-4-H6	684 bp	4	227 aa	4	1d 2m	V F M S	> xenopus EST > repetitive element [CDS CATEGORY]
S10-4-H7	692 bp	1	230 aa	2	2p 1d 2m	V F M S	> exostosis 2 (partial) > Duplicate [CDS CATEGORY]
S10-4-H8	653 bp	5	217 aa	5	1p 1d 2m	V F M S	> cdc23 (FL) [CDS CATEGORY]
S10-4-H9	667 bp	4	222 aa	4	1p 1d 2m	V F M S	> RAC-2/RhoG > Duplicate [CDS CATEGORY]

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Questions and comments on **MAGPIE** are welcome! Send to [gaasterland@rockefeller.edu](mailto:gaasterland@rockefeller.edu) [Terry Gaasterland] or [sensencw@niji.imb.nrc.ca](mailto:sensencw@niji.imb.nrc.ca)

TOE220 2460T660

# M A G P I E - P R O J E C T - xenopus

GROUP: S10-5

STATE: protein\_dna

<S10-4 -- S10-6 >  
Mon Jul 10 08:50:36 EDT 2000

sort by signature

STATUS | CONTIG INFO | ANNOTATIONS | PRIMERS | ANNOTATOR

The S10-5 group contains 96 contigs.

ID	Bases	N	AA	X	Evidence	Forms	EC	Description	Oth Rep
S10-5-A1	714 bp	99	237 aa	38	1p 1d 2m	V F M S		> ODC (partial) > duplicate [CDS CATEGORY]	
S10-5-A10	696 bp	1	231 aa	1	2p 1d 2m	V F M S		> Ribonuclease H1 (FL) > empty description [CDS CATEGORY]	
S10-5-A11	683 bp	8	227 aa	7	1p 1d 2m	V F M S		> HSP-70 (FL) [CDS CATEGORY]	
S10-5-A12	685 bp	3	228 aa	2	1p 1d 2m	V F M S		> Inner Centromer Protein (FL) [CDS CATEGORY]	
S10-5-A2	785 bp	10	261 aa	9	2p 1d 2m	V F M S		> Potassium channel ORF (FL) [CDS CATEGORY]	
S10-5-A3	705 bp	4	234 aa	4	2p 1d 2m	V F M S		> Rab/c-mel gtp-binding protein (FL) [CDS CATEGORY]	
S10-5-A4	720 bp	6	239 aa	6	1d 2m	V F M S		> Xenopus EST	
S10-5-A5	717 bp	3	238 aa	3	1p 1d 2m	V F M S		> Cyt c oxidase subunit I (partial) > Duplicate [CDS Catego	
S10-5-A6	717 bp	4	238 aa	3	1p 1d 2m	V F M S		> cyclin G2 (FL) [CDS CATEGORY]	
S10-5-A7	717 bp	1	238 aa	1	1p 1d 2m	V F M S		> PROTEIN TRANSLATION FACTOR SUI1 [CDS CATEGORY]	
S10-5-A8	691 bp	65	230 aa	51	3p 1d 2m	V F M S		> RRM RNA binding protein gry-rbp (FL) > Duplicate [CDS CAT	
S10-5-A9	678 bp	1	225 aa	1	1p 1d 2m	V F M S		> H3 histone, H3.3 (FL) > Duplicate [CDS CATEGORY]	
S10-5-B1	776 bp	7	258 aa	8	1d 2m	V F M S		> Xenopus EST	
S10-5-B10	694 bp	3	231 aa	4	1p 1d 2m	V F M S		> cold-inducible rna-binding protein (partial) > duplicate	

S10-5-B11	734 bp	4	244 aa	4	1p 1d 2m	V F M S	> tftis elongation factor (FL) > duplicate [CDS CATEGORY]
S10-5-B12	682 bp	1	227 aa	2	1p 1d 2m	V F M S	> XFD-12 [CDS CATEGORY]
S10-5-B2	737 bp	1	245 aa	2	2p 1d 2m	V F M S	> RING finger protein (partial) [CDS CATEGORY]
S10-5-B3	712 bp	3	237 aa	2	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-5-B4	718 bp	2	239 aa	2	1p 1d 2m	V F M S	> ubiquinol-cytochrome C reductase [CDS CATEGORY]
S10-5-B5	714 bp	2	237 aa	1	3p 1d 2m	V F M S	> Similar to Ribosomal L14 protein (FL) [CDS CATEGORY]
S10-5-B6	717 bp	2	238 aa	1	1p 1d 2m	V F M S	> G2/Mitotic Specific cyclin B1' (FL) [CDS CATEGORY]
S10-5-B7	710 bp	1	236 aa	2	1d 2m	V F M S	> xenopus EST > Duplicate [CDS CATEGORY]
S10-5-B8	678 bp	6	225 aa	6	2p 1d 2m	V F M S	> Flash (FL) [CDS CATEGORY]
S10-5-B9	695 bp	8	231 aa	6	1p 1d 2m	V F M S	> Oct-1 (FL) [CDS CATEGORY]
S10-5-C1	714 bp	3	237 aa	3	1p 1d 2m	V F M S	> kina1038 like (partial) > similar to EF1a (partial) [CDS
S10-5-C10	667 bp	1	222 aa	2	3p 1d 2m	V F M S	> xenopus EST > duplicate [CDS CATEGORY]
S10-5-C11	166 bp	19	55 aa	15	1d	V F M S	> Xenopus EST
S10-5-C12	733 bp	7	244 aa	5	2p 1d 2m	V F M S	> ATPase, Cu++ transporting (partial) [CDS CATEGORY]
S10-5-C2	690 bp	9	229 aa	6	3p 1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-5-C3	726 bp	47	241 aa	34	3p 1d 2m	V F M S	> cyclin A1 (FL) [CDS CATEGORY]
S10-5-C4	720 bp	8	239 aa	7	1p 1d 2m	V F M S	> PLX-1 Polo like kinase (partial) [CDS CATEGORY]
S10-5-C5	731 bp	5	243 aa	5	1p 1d 2m	V F M S	> RanBPM centrosomal protein (FL) [CDS CATEGORY]
S10-5-C6	733 bp	8	244 aa	6	1p 1d 2m	V F M S	> Arg/Ser-rich 5 splicing factor (FL) > duplicate [CDS CATE
S10-5-C7	715 bp	3	238 aa	4	2p 1d 2m	V F M S	> BMP-7 (partial) [CDS CATEGORY]
S10-5-C8	714 bp	8	237 aa	6	1d 2m	V F M S	> Cyclin A2 (FL) [CDS CATEGORY]
S10-5-C9	736 bp	4	245 aa	4	1p 1d 2m	V F M S	> t-complex protein 1, epsilon subunit (FL) [CDS CATEGORY]
S10-5-D1	737 bp	11	245 aa	10	3p 1d 2m	V F M S	> Histone deacetylase (partial) [CDS CATEGORY]
S10-5-D10	667 bp	1	222 aa	2	1p 1d 2m	V F M S	> ankyrin repeat G9A (partial) [CDS CATEGORY]
S10-5-D11	667 bp	2	222 aa	2	1p 1d 2m	V F M S	> prolyl 4-hydroxylase (FL) [CDS CATEGORY]
S10-5-D12	685 bp	1	228 aa	2	1p 1d 2m	V F M S	> rab interacting kinesin (partial) [CDS CATEGORY]
S10-5-D2	706 bp	5	235 aa	5	2p 1d 2m	V F M S	> C elegans ORF (FL) > Xenopus EST [CDS CATEGORY]
S10-5-D3	724 bp	7	241 aa	6	2p 1d 2m	V F M S	> Sno notch pathway component (partial) [CDS CATEGORY]
S10-5-D4	715 bp	1	238 aa	2	2p 1d 2m	V F M S	> glutamic acid-rich protein precursor (FL) > Xenopus EST [
S10-5-D5	729 bp	6	242 aa	5	3p 1d 2m	V F M S	> SP3 like (FL) [CDS CATEGORY]

TABLE 4: 2460T.660

S10-5-D6	717 bp	2	238 aa		1p 1d 2m	V F M S
S10-5-D7	482 bp	1	160 aa	2	3p 1d 2m	V F M S
S10-5-D8	714 bp	5	237 aa	5	1p 1d 2m	V F M S
S10-5-D9	676 bp	2	225 aa	3	1p 1d 2m	V F M S
S10-5-E1	734 bp	3	244 aa	3	3p 1d 2m	V F M S
S10-5-E10	687 bp	2	228 aa	2	1d 2m	V F M S
S10-5-E11	829 bp	242	276 aa	155	1d 2m	V F M S
S10-5-E12	683 bp	6	227 aa	5	1d 2m	V F M S
S10-5-E2	737 bp	5	245 aa	5	1p 1d 2m	V F M S
S10-5-E3	722 bp	3	240 aa	3	1p 1d 2m	V F M S
S10-5-E4	713 bp	1	237 aa	2	2p 1d 2m	V F M S
S10-5-E5	718 bp	6	239 aa	6	3p 1d 2m	V F M S
S10-5-E6	718 bp	58	239 aa	56	1d 2m	V F M S
S10-5-E7	714 bp	1	237 aa	1	1p 1d 2m	V F M S
S10-5-E8	572 bp	1	190 aa	2	1d 2m	V F M S
S10-5-E9	694 bp	4	231 aa	5	1d 2m	V F M S
S10-5-F1	787 bp	3	262 aa	3	1p 1d 2m	V F M S
S10-5-F10	738 bp	4	245 aa	3	1d 2m	V F M S
S10-5-F11	694 bp	19	231 aa	17	1d 2m	V F M S
S10-5-F12	682 bp	5	227 aa	6	1d 2m	V F M S
S10-5-F2	770 bp	105	256 aa	84	2d 2m	V F M S
S10-5-F3	549 bp	3	182 aa	2	1p 1d 2m	V F M S
S10-5-F4	734 bp	20	244 aa	13	1p 1d 2m	V F M S
S10-5-F5	730 bp	5	243 aa	4	3p 1d 2m	V F M S
S10-5-F6	715 bp	7	238 aa	7	3p 1d 2m	V F M S
S10-5-F7	716 bp	1	238 aa	1	1p 1d 2m	V F M S
S10-5-F8	712 bp	15	237 aa	12	3p 1d 2m	V F M S
S10-5-F9	480 bp	11	159 aa	7	2p 1d 2m	V F M S
S10-5-G1	735 bp	1	244 aa		1d 2m	V F M S

> poly binding protein 2 (partial) > duplicate [CDS CATEGOR
> Xenopus EST [CDS CATEGORY]
> Ser/Thr phosphatase pp2a-4 (FL) > Duplicate [CDS CATEGORY
> RAB6 interacting, kinesin-like (FL) [CDS CATEGORY]
> Xenopus EST > empty description [CDS CATEGORY]
> Xenopus EST > Repetitive Sequence [CDS CATEGORY]
> Xenopus EST
> xElr-C elav-like (FL) [CDS CATEGORY]
> JNK protein kinase (partial) [CDS CATEGORY]
> NifU like (FL) [CDS CATEGORY]
> CGI-73 conserved protein (FL) [CDS CATEGORY]
> Vector > Duplicate [CDS CATEGORY]
> Vector > duplicate [CDS CATEGORY]
> peroxisomal ca-dependent solute carrier (FL) [CDS CATEGOR
> Xenopus EST
> Xenopus EST [CDS CATEGORY]
> U1 snrnp A PROTEIN (FL) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> Xenopus EST
> Xenopus EST [CDS CATEGORY]
> Xenopus EST
> CAF-1 P55 subunit/ RB binding Prot 7(partial) [CDS CATEGO
> BC-2 [CDS CATEGORY]
> splicing factor > xenopus EST [CDS CATEGORY]
> Vector > Duplicate [CDS CATEGORY]
> matrin 3 (FL) [CDS CATEGORY]
> Xenopus EST > tudor repeat associator with PCTAIRE (partia
> ferritin H (FL) > duplicate [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]



TABLE 2

S10-5-G10	689 bp	2	229 aa	3	1p 1d 2m	V F M S	> EF1-ALPHA (FL) [CDS CATEGORY]
S10-5-G11	665 bp	3	221 aa	4	1p 1d 2m	V F M S	> ATP synthase beta-subunit (FL) > Duplicate [CDS CATEGORY]
S10-5-G12	670 bp	7	223 aa	8	1p 1d 2m	V F M S	> splicing factor (FL) [CDS CATEGORY]
S10-5-G2	864 bp	215	287 aa	152	1d 2m	V F M S	> Xenopus EST
S10-5-G3	696 bp	7	231 aa	6	1p 1d 2m	V F M S	> protein phosphatase 5 (FL) [CDS CATEGORY]
S10-5-G4	745 bp	74	248 aa	56	1d 2m	V F M S	> Xenopus EST
S10-5-G5	723 bp	1	240 aa	1	2p 1d 2m	V F M S	> sec61 alpha subunit (partial) > vector [CDS CATEGORY]
S10-5-G6	722 bp	8	240 aa	8	1d 2m	V F M S	> Xenopus EST
S10-5-G7	701 bp	3	233 aa	3	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-5-G8	711 bp	1	236 aa	1	1p 1d 2m	V F M S	> vacuolar sorting protein VPS29 (FL) [CDS CATEGORY]
S10-5-G9	691 bp	1	230 aa	1	3p 1d 2m	V F M S	> Xenopus EST > Fat Facets 3' homology [CDS CATEGORY]
S10-5-H1	778 bp	11	259 aa	11	1p 1d 2m	V F M S	> cdc-6 (FL) [CDS CATEGORY]
S10-5-H10	691 bp	3	230 aa	4	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-5-H11	689 bp	8	229 aa	7	1p 1d 2m	V F M S	> mapk-activated protein kinase 2 (FL) > Duplicate [CDS CAT
S10-5-H12	683 bp	8	227 aa	7	1p 1d 2m	V F M S	> casein kinase II, alpha' (partial) [CDS CATEGORY]
S10-5-H2	715 bp	194	238 aa	67	1d 2m	V F M S	> Xenopus EST
S10-5-H3	720 bp	4	239 aa	4	1p 1d 2m	V F M S	> cyclin A1 (partial) [CDS CATEGORY]
S10-5-H4	726 bp	26	241 aa	21	2p 1d 2m	V F M S	> cytochrome c oxidase subunit I (M10217) > Vector [CDS CAT
S10-5-H5	726 bp	5	241 aa	4	2p 1d 2m	V F M S	> Asparaginase similarity (nFL) [CDS CATEGORY]
S10-5-H6	716 bp	6	238 aa	5	1p 1d 2m	V F M S	> dihydrolipoamide succinyltransferase (FL) [CDS CATEGORY]
S10-5-H7	815 bp	164	271 aa	120	1d 2m	V F M S	> Xenopus EST
S10-5-H8	702 bp	6	233 aa	6	1p 1d 2m	V F M S	> tripeptidylpeptidase II (nFL) [CDS CATEGORY]
S10-5-H9	697 bp	8	232 aa	9	2p 1d 2m	V F M S	> sorting nexin 1 (partial) [CDS CATEGORY]

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Questions and comments on MAGPIE are welcome! Send to [gaasterland@rockefeller.edu](mailto:gaasterland@rockefeller.edu) [Terry Gaasterland] or [sensencw@niji.imb.nrc.ca](mailto:sensencw@niji.imb.nrc.ca)

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# MAGPIE-PROJECT - xenopus

GROUP: S10-6

STATE: protein\_dna

&lt;S10-5 -- S10-7 &gt;

Mon Jul 10 08:50:36 EDT 2000

sort by signature

STATUS | CONTIG INFO | ANNOTATIONS | PRIMERS | ANNOTATOR

The S10-6 group contains 96 contigs.

ID	Bases	N	AA	X	Evidence	Forms	EC	Description	Oth Repo
S10-6-A1	682 bp	1	227 aa	1	1d 2m	V F M S		> Xenopus EST [CDS CATEGORY]	
S10-6-A10	696 bp	2	231 aa	2	1p 1d 2m	V F M S		> hmg-1 (partial) > duplicate [CDS CATEGORY]	
S10-6-A11	681 bp		226 aa		1d 2m	V F M S		> Xenopus EST	
S10-6-A12	740 bp	103	246 aa	83	1d 2m	V F M S		> Xenopus EST	
S10-6-A2	652 bp	1	217 aa	2	1d 2m	V F M S		> Xenopus EST	
S10-6-A3	668 bp	4	222 aa	4	1p 1d 2m	V F M S		> RING finger protein (FL) [CDS CATEGORY]	
S10-6-A4	715 bp	125	238 aa	70	1d 2m	V F M S		> Xenopus EST	
S10-6-A5	710 bp	5	236 aa	4	1p 1d 2m	V F M S		> polyA binding protein II (FL) [CDS CATEGORY]	
S10-6-A6	711 bp	3	236 aa	2	1p 1d 2m	V F M S		> Arg/Ser rich splicing factor 11 (FL) [CDS CATEGORY]	
S10-6-A7	695 bp	1	231 aa	2	1p 1d 2m	V F M S		> xOct-25 (FL) [CDS CATEGORY]	
S10-6-A8	696 bp	1	231 aa	1	3p 1d 2m	V F M S		> xNF-7 (FL) [CDS CATEGORY]	
S10-6-A9	747 bp	71	248 aa	51	1p 1d 2m	V F M S		> ODC (FL) > Duplicate [CDS CATEGORY]	
S10-6-B1	679 bp	2	226 aa	3	1d 2m	V F M S		> Xenopus EST	
S10-6-B10	695 bp	1	231 aa	2	1p 1d 2m	V F M S		> cdc2 kinase (FL) [CDS CATEGORY]	

[illegible]

S10-6-B11	682 bp		227 aa	1	1p 1d 2m	V F M S	> RAC Protein Kinase BETA (FL) [CDS CATEGORY]
S10-6-B12	659 bp	2	219 aa	3	3p 1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-6-B2	663 bp		220 aa		1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-6-B3	672 bp	1	223 aa	1	2p 1d 2m	V F M S	> kiaa0948 protein (FL) > Xenopus EST [CDS CATEGORY]
S10-6-B4	736 bp	171	245 aa	111	1d 2m	V F M S	> Xenopus EST
S10-6-B5	771 bp	16	256 aa	15	1p 1d 2m	V F M S	> lamina associated protein 2-beta isoform (FL) [CDS CATEGORY]
S10-6-B6	726 bp	62	241 aa	47	3p 1d 2m	V F M S	> cdc2/cdk2,4-Activating kinase (nFL) [CDS CATEGORY]
S10-6-B7	750 bp	3	249 aa	3	3p 1d 2m	V F M S	> Syntaxin-11 like (FL) [CDS CATEGORY]
S10-6-B8	529 bp	3	176 aa	3	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-6-B9	718 bp	4	239 aa	4	1d 2m	V F M S	> Xenopus EST
S10-6-C1	678 bp		225 aa		1p 1d 2m	V F M S	> transcriptional regulator protein (FL) [CDS CATEGORY]
S10-6-C10	689 bp	2	229 aa	2	1p 1d 2m	V F M S	> phosphatidylinositol glycan, class C (partial) [CDS CATEGORY]
S10-6-C11	680 bp		226 aa	1	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-6-C12	680 bp		226 aa	1	1d 2m	V F M S	> retrotransposon element [CDS CATEGORY]
S10-6-C2	679 bp		226 aa	1	3p 1d 2m	V F M S	> Desmoplakin I (partial) [CDS CATEGORY]
S10-6-C3	676 bp	1	225 aa	2	1p 1d 2m	V F M S	> polyubiquitin (partial) [CDS CATEGORY]
S10-6-C4	672 bp	2	223 aa	2	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-6-C5	766 bp	8	255 aa	7	2p 1d 2m	V F M S	> opioid growth factor receptor (partial) [CDS CATEGORY]
S10-6-C6	710 bp	2	236 aa	3	2p 1d 2m	V F M S	> growth arrest specific 11 (partial) [CDS CATEGORY]
S10-6-C7	697 bp	1	232 aa	2	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-6-C8	656 bp	1	218 aa	1	1p 1d 2m	V F M S	> disulfide-isomerase xER60 [CDS CATEGORY]
S10-6-C9	315 bp	42	104 aa	27	1d	V F M S	> Nuc pore complex assoc (partial) [CDS CATEGORY]
S10-6-D1	678 bp		225 aa		1d 2m	V F M S	> XFG 5-1/ XFG 5-2 > duplicate [CDS CATEGORY]
S10-6-D10	693 bp	2	230 aa	2	1p 1d 2m	V F M S	> Arg/Ser-rich 5 splicing factor (FL) > duplicate [CDS CATEGORY]
S10-6-D11	681 bp		226 aa		1p 1d 2m	V F M S	> PBK1 protein (nFL) [CDS CATEGORY]
S10-6-D12	666 bp		221 aa		3p 1d 2m	V F M S	> Xenopus EST RT homology [CDS CATEGORY]
S10-6-D2	677 bp	1	225 aa	2	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-6-D3	670 bp	1	223 aa	2	2p 1d 2m	V F M S	> thyroid hormone receptor coactivating protein (partial) [
S10-6-D4	645 bp	2	214 aa	1	3p 1d 2m	V F M S	> Xenopus EST > Wiskott-Aldrich syndrome interacting protein
S10-6-D5	706 bp	6	235 aa	7	1p 1d 2m	V F M S	> cold-inducible rna-binding protein (partial) > duplicate



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S10-6-D6	706 bp	2	235 aa	3	1d 2m	V F M S
S10-6-D7	737 bp	116	245 aa	77	1d 2m	V F M S
S10-6-D8	759 bp	15	252 aa	7	1d 2m	V F M S
S10-6-D9	695 bp	4	231 aa	4	1d 2m	V F M S
S10-6-E1	667 bp	34	222 aa	14	3p 1d 2m	V F M S
S10-6-E10	686 bp	1	228 aa	2	1p 1d 2m	V F M S
S10-6-E11	660 bp	17	219 aa	11	1p 1d 2m	V F M S
S10-6-E12	680 bp	2	226 aa	2	3p 1d 2m	V F M S
S10-6-E2	678 bp	1	225 aa	1	1p 1d 2m	V F M S
S10-6-E3	724 bp	10	241 aa	8	2p 1d 2m	V F M S
S10-6-E4	713 bp	7	237 aa	5	1p 1d 2m	V F M S
S10-6-E5	667 bp	9	222 aa	8	1p 1d 2m	V F M S
S10-6-E6	758 bp	7	252 aa	7	1p 1d 2m	V F M S
S10-6-E7	692 bp	2	230 aa	2	1p 1d 2m	V F M S
S10-6-E8	655 bp	2	218 aa	2	1p 1d 2m	V F M S
S10-6-E9	695 bp	3	231 aa	4	1p 1d 2m	V F M S
S10-6-F1	756 bp	11	251 aa	6	3p 1d 2m	V F M S
S10-6-F10	688 bp	1	229 aa	2	3p 1d 2m	V F M S
S10-6-F11	683 bp	1	227 aa	1	1d 2m	V F M S
S10-6-F12	668 bp		222 aa	1	1p 1d 2m	V F M S
S10-6-F2	702 bp	120	233 aa	90	1d 2m	V F M S
S10-6-F3	652 bp	4	217 aa	4	1p 1d 2m	V F M S
S10-6-F4	672 bp	3	223 aa	3	1d 2m	V F M S
S10-6-F5	757 bp	2	252 aa	1	1p 1d 2m	V F M S
S10-6-F6	708 bp	7	235 aa	6	1p 1d 2m	V F M S
S10-6-F7	668 bp	1	222 aa	2	1p 1d 2m	V F M S
S10-6-F8	691 bp	1	230 aa	2	1d 2m	V F M S
S10-6-F9	673 bp	4	224 aa	4	1d 2m	V F M S
S10-6-G1	661 bp	1	220 aa	1	1p 1d 2m	V F M S

> Xenopus repetitive sequence
> Xenopus
> Xenopus EST [CDS CATEGORY]
> Xenopus EST > Duplicate [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> phosphoprotein phosphatase (FL) [CDS CATEGORY]
> alternative splicing factor asf-3 (FL) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> EF1-alpha (partial) [CDS CATEGORY]
> p47 (FL) > empty description [CDS CATEGORY]
> pre-mRNase cleavage factor Im (FL) [CDS CATEGORY]
> protein phosphatase 2C beta (FL) > Duplicate [CDS CATEGORY]
> hmg-1 (partial) > duplicate [CDS CATEGORY]
> histone H3.3 (partial) [CDS CATEGORY]
> monocarboxylate transporter 1 (FL) [CDS CATEGORY]
> Histone-Binding Protein N1/N2 (FL) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> pyruvate KINASE, muscle isozyme (partial) [CDS CATEGORY]
> Xenopus EST
> aldo-keto reductase (FL) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> biphenyl hydrolase-related protein (FL) [CDS CATEGORY]
> Keratin, Type II cytoskeletal 8 (FL) > Duplicate [CDS CATEGORY]
> cdc47-2p (FL) > duplicate [CDS CATEGORY]
> Xenopus EST
> Xenopus EST > Duplicate [CDS CATEGORY]
> kiao0370 (FL) [CDS CATEGORY]

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S10-6-G10	688 bp	2	229 aa	3	2p 1d 2m	V F M S	> kelch MIPP like (partial) [CDS CATEGORY]
S10-6-G11	673 bp	6	224 aa	5	1p 1d 2m	V F M S	> claudin-7 (FL) > Duplicate [CDS CATEGORY]
S10-6-G12	680 bp	3	226 aa	4	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-6-G2	674 bp	5	224 aa	5	1d 2m	V F M S	> xenopus EST > Duplicate [CDS CATEGORY]
S10-6-G3	743 bp	15	247 aa	10	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-6-G4	658 bp	1	219 aa	2	1p 1d 2m	V F M S	> XlhnRNPL (partial) [CDS CATEGORY]
S10-6-G5	756 bp	13	251 aa	11	1d 2m	V F M S	> Xenopus EST
S10-6-G6	715 bp	3	238 aa	4	1p 1d 2m	V F M S	> DNA replication factor mcm6b (partial) [CDS CATEGORY]
S10-6-G7	744 bp	5	247 aa	3	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-6-G8	670 bp	1	223 aa	2	1p 1d 2m	V F M S	> eIF2 gamma (FL) [CDS CATEGORY]
S10-6-G9	690 bp	1	229 aa	1	1p 1d 2m	V F M S	> protein disulfide isomerase-related xERP-72 (partial) [CD
S10-6-H1	675 bp	1	224 aa	1	3p 1d 2m	V F M S	> Zn Finger protein (partial) [CDS CATEGORY]
S10-6-H10	655 bp	1	218 aa	2	1p 1d 2m	V F M S	> 40S ribosomal protein S4 (nFL) > Duplicate [CDS CATEGORY]
S10-6-H11	688 bp	5	229 aa	5	3p 1d 2m	V F M S	> 5-Aminolevulinic acid Synthase (FL) [CDS CATEGORY]
S10-6-H12	681 bp	1	226 aa	1	1d 2m	V F M S	> Xenopus EST
S10-6-H2	677 bp	9	225 aa	7	3p 1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-6-H3	657 bp		218 aa		1d 2m	V F M S	> Nucleolin (partial) [CDS CATEGORY]
S10-6-H4	717 bp	4	238 aa	3	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-6-H5	761 bp	11	253 aa	8	1p 1d 2m	V F M S	> hnrap A2/B1 (FL) > Duplicate [CDS CATEGORY]
S10-6-H6	709 bp	5	236 aa	4	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-6-H7	750 bp	7	249 aa	7	1p 1d 2m	V F M S	> Keratin, Type II cytoskeletal 8 (FL) > Duplicate [CDS CAT
S10-6-H8	738 bp	3	245 aa	3	2p 1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-6-H9	687 bp	3	228 aa	3	2p 1d 2m	V F M S	> CGI-86 like (FL) [CDS CATEGORY]

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Questions and comments on MAGPIE are welcome! Send to [gaasterland@rockefeller.edu](mailto:gaasterland@rockefeller.edu) [Terry Gaasterland] or [sensencw@niji.imb.mrc.ca](mailto:sensencw@niji.imb.mrc.ca)

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## MAGPIE - PROJECT - xenopus

GROUP: S10-7

STATE: protein\_dna

<S10-6 -- S10-8 >

Mon Jul 10 08:50:36 EDT 2000

sort by signature

STATUS | CONTIG INFO | ANNOTATIONS | PRIMERS | ANNOTATOR

The S10-7 group contains 96 contigs.

ID	Bases	N	AA	X	Evidence	Forms	EC	Description	Oth Repo
S10-7-A1	743 bp	4	247 aa	5	1p 1d 2m	V F M S		> glycerol kinase (FL) [CDS CATEGORY]	
S10-7-A10	675 bp	5	224 aa	4	1p 1d 2m	V F M S		> 5'-nucleotidase (FL) [CDS CATEGORY]	
S10-7-A11	670 bp	13	223 aa	14	1d 2m	V F M S		> Xenopus EST [CDS CATEGORY]	
S10-7-A12	706 bp	3	235 aa	3	1d 2m	V F M S		> Xenopus EST [CDS CATEGORY]	
S10-7-A2	728 bp	3	242 aa	4	1p 1d 2m	V F M S		> 3-oxoacid CoA transferase (FL) [CDS CATEGORY]	
S10-7-A3	789 bp	20	262 aa	15	2p 1d 2m	V F M S		> U1 snRNP 70 KD (FL) [CDS CATEGORY]	
S10-7-A4	766 bp	3	255 aa	4	1p 1d 2m	V F M S		> cytochrome B (FL) [CDS CATEGORY]	
S10-7-A5	800 bp	8	266 aa	6	1p 1d 2m	V F M S		> Suppressor of hairless protein 1 (FL) [CDS CATEGORY]	
S10-7-A6	725 bp	5	241 aa	5	1p 1d 2m	V F M S		> protein tyrosine kinase 9 (FL) > Xenopus EST [CDS CATEGORY]	
S10-7-A7	716 bp	2	238 aa	2	1d 2m	V F M S		> Xenopus EST [CDS CATEGORY]	
S10-7-A8	663 bp	4	220 aa	4	1p 1d 2m	V F M S		> TIAM/SIF (partial) [CDS CATEGORY]	
S10-7-A9	761 bp	18	253 aa	16	1p 1d 2m	V F M S		> ubiquitin-like protein smt3a (FL) > Duplicate [CDS CATEGORY]	
S10-7-B1	736 bp	2	245 aa	2	1p 1d 2m	V F M S		> DG42 protein (FL) [CDS CATEGORY]	
S10-7-B10	709 bp	1	236 aa	2	1p 1d	V F M S		> cytochrome c oxidase subunit I (nFL) > Duplicate [CDS CATEGORY]	

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S10-7-B11	676 bp	3	225 aa	3	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-7-B12	699 bp	4	232 aa	4	1p 1d 2m	V F M S	> nice-3 protein (FL) [CDS CATEGORY]
S10-7-B2	790 bp	9	263 aa	8	3p 1d 2m	V F M S	> KIAA0886 protein (partial) [CDS CATEGORY]
S10-7-B3	777 bp	6	258 aa	6	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-7-B4	724 bp	3	241 aa	2	1p 1d 2m	V F M S	> Xenopus EST > Human hypothetical (partial) [CDS CATEGORY]
S10-7-B5	712 bp	5	237 aa	5	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-7-B6	718 bp	3	239 aa	2	3p 1d 2m	V F M S	> Cyclin 1 (FL) [CDS CATEGORY]
S10-7-B7	709 bp	8	236 aa	7	1d 2m	V F M S	> Xenopus EST > Repetitive element [CDS CATEGORY]
S10-7-B8	716 bp	4	238 aa	5	1p 1d 2m	V F M S	> Human hypothetical (FL) [CDS CATEGORY]
S10-7-B9	709 bp	2	236 aa	3	1p 1d 2m	V F M S	> tyrosine-protein kinase src-2 (P60-SRC-2) (FL) [CDS CATEG
S10-7-C1	780 bp	6	259 aa	5	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-7-C10	764 bp	8	254 aa	7	1p 1d 2m	V F M S	> sperm surface protein/JNK/SAPK interacting (partial) [CDS
S10-7-C11	699 bp	5	232 aa	5	1p 1d 2m	V F M S	> k1aa0421/lambda/iota protein kinase C (partial) [CDS CATE
S10-7-C12	694 bp	1	231 aa	2	1p 1d 2m	V F M S	> Variant Histone H2A.Z12 (FL) [CDS CATEGORY]
S10-7-C2	780 bp	9	259 aa	8	1p 1d 2m	V F M S	> ADOMETDC (FL) [CDS CATEGORY]
S10-7-C3	791 bp	7	263 aa	6	3p 1d 2m	V F M S	> thioredoxin interacting factor (partial) [CDS CATEGORY]
S10-7-C4	698 bp	1	232 aa	2	3p 1d 2m	V F M S	> Xenopus EST > C elegans ORF (FL) [CDS CATEGORY]
S10-7-C5	707 bp	2	235 aa	3	1p 1d 2m	V F M S	> hepatoma-derived growth factor (FL) [CDS CATEGORY]
S10-7-C6	723 bp	3	240 aa	2	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-7-C7	716 bp	4	238 aa	4	1p 1d 2m	V F M S	> BMP-7 (FL) [CDS CATEGORY]
S10-7-C8	712 bp	4	237 aa	5	1p 1d 2m	V F M S	> claudin-7 (FL) > Duplicate [CDS CATEGORY]
S10-7-C9	769 bp	7	256 aa	6	1p 1d 2m	V F M S	> G protein pathway suppressor 1 (FL) [CDS CATEGORY]
S10-7-D1	732 bp	5	243 aa	5	1d 2m	V F M S	> Xenopus EST
S10-7-D10	703 bp	5	234 aa	4	1p 1d 2m	V F M S	> fatvg (FL) [CDS CATEGORY]
S10-7-D11	683 bp	1	227 aa	2	1p 1d 2m	V F M S	> Cyt c oxidase subunit I (partial) [CDS CATEGORY]
S10-7-D12	702 bp	42	233 aa	18	1d	V F M S	> mitochondria > Similar [CDS CATEGORY]
S10-7-D2	751 bp	21	250 aa	18	1p 1d 2m	V F M S	> protein kinase, cAMP-dependent, regulatory (partial) [CDS
S10-7-D3	596 bp	10	198 aa	10	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-7-D4	788 bp	11	262 aa	10	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-7-D5	683 bp	3	227 aa	3	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]

S10-7-D6	717 bp	1	238 aa	1	1p 1d 2m	V F M S
S10-7-D7	779 bp	4	259 aa	5	1d 2m	V F M S
S10-7-D8	763 bp	12	254 aa	9	1d 2m	V F M S
S10-7-D9	756 bp	8	251 aa	7	1d 2m	V F M S
S10-7-E1	734 bp	1	244 aa	2	1d 2m	V F M S
S10-7-E10	714 bp	3	237 aa	3	1p 1d 2m	V F M S
S10-7-E11	684 bp	4	227 aa	4	1d 2m	V F M S
S10-7-E12	642 bp		213 aa		1p 1d 2m	V F M S
S10-7-E2	736 bp	4	245 aa	5	1p 1d 2m	V F M S
S10-7-E3	712 bp	3	237 aa	4	3p 1d 2m	V F M S
S10-7-E4	275 bp	3	91 aa	3	1d 2m	V F M S
S10-7-E5	718 bp	1	239 aa	2	2p 1d 2m	V F M S
S10-7-E6	707 bp	1	235 aa	2	2p 1d 2m	V F M S
S10-7-E7	710 bp	9	236 aa	9	1d	V F M S
S10-7-E8	696 bp	50	231 aa	35	3p 1d 2m	V F M S
S10-7-E9	771 bp	17	256 aa	15	1p 1d 2m	V F M S
S10-7-F1	734 bp	5	244 aa	5	1p 1d	V F M S
S10-7-F10	711 bp	5	236 aa	4	1p 1d 2m	V F M S
S10-7-F11	753 bp	3	250 aa	3	1p 1d 2m	V F M S
S10-7-F12	367 bp	6	122 aa	6	1d 2m	V F M S
S10-7-F2	720 bp	5	239 aa	5	3p 1d 2m	V F M S
S10-7-F3	727 bp	5	242 aa	4	1d 2m	V F M S
S10-7-F4	723 bp	1	240 aa	1	1p 1d 2m	V F M S
S10-7-F5	725 bp	5	241 aa	4	1d 2m	V F M S
S10-7-F6	795 bp	11	264 aa	8	1p 1d 2m	V F M S
S10-7-F7	715 bp	2	238 aa	3	3p 1d 2m	V F M S
S10-7-F8	715 bp	2	238 aa	3	1d 2m	V F M S
S10-7-F9	720 bp	9	239 aa	7	1d 2m	V F M S
S10-7-G1	735 bp	5	244 aa	5	1p 1d 2m	V F M S

> phospholipase C, epsilon (partial) [CDS CATEGORY]
> Xenopus EST
> xenopus EST > duplicate [CDS CATEGORY]
> Xenopus EST
> Xenopus EST [CDS CATEGORY]
> p75-like transmembrane protein fullback (FL) [CDS CATEGORY]
> Xenopus EST
> hn ribonucleoprotein R (partial) [CDS CATEGORY]
> ADP-ribosylation factor 4 (FL) [CDS CATEGORY]
> Xenopus EST > Synaptic vesicle Prot. Vst-1 (FL) [CDS CATEGORY]
> Xenopus EST > Conserved Element [CDS CATEGORY]
> Xenopus EST > Transposase (minus Strand) [CDS CATEGORY]
> Hypothetical Protein (nFL) [CDS CATEGORY]
> xenopus EST > Duplicate [CDS CATEGORY]
> Vector > Duplicate [CDS CATEGORY]
> Pak-2 (partial) [CDS CATEGORY]
> Protein kinase FNK/SNK (partial?) [CDS CATEGORY]
> mitochondrial uncoupling protein 2 (partial) [CDS CATEGORY]
> tyrosine kinase JAK1 (partial) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> xMek-3 (FL) [CDS CATEGORY]
> Xenopus EST > Conserved element [CDS CATEGORY]
> Paraneoplastic antigen (partial) [CDS CATEGORY]
> Xenopus EST
> calumenin (FL) [CDS CATEGORY]
> cdc25C2 (partial) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> G2/MITOTIC-SPECIFIC cyclin B1 (FL) > Duplicate [CDS CATEGORY]
> NADH-cytochrome B5 reductase (FL) [CDS CATEGORY]



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S10-7-G10	703 bp	6	234 aa	5	1p 1d 2m	V F M S	> amino acid transporter B0+ (partial) [CDS CATEGORY]
S10-7-G11	752 bp	4	250 aa	4	1p 1d 2m	V F M S	> FAT-3 alcohol dehydrogenase like (FL) [CDS CATEGORY]
S10-7-G12	706 bp	7	235 aa	7	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-7-G2	738 bp	14	245 aa	12	2p 1d 2m	V F M S	> translation repressor NAT1 (partial) [CDS CATEGORY]
S10-7-G3	709 bp	7	236 aa	7	1d 2m	V F M S	> Xenopus EST
S10-7-G4	736 bp	5	245 aa	6	1p 1d 2m	V F M S	> fls353 (partial) [CDS CATEGORY]
S10-7-G5	719 bp	2	239 aa	3	2p 1d 2m	V F M S	> k1aa1095 protein (FL) > Xenopus EST [CDS CATEGORY]
S10-7-G6	592 bp	11	197 aa	7	1d 2m	V F M S	> Xenopus EST
S10-7-G7	719 bp	2	239 aa	1	2p 1d 2m	V F M S	> U3 snoRNP-associated 55-kd protein (FL) > Xenopus EST [CD
S10-7-G8	711 bp		236 aa		1p 1d 2m	V F M S	> G9A (partial) [CDS CATEGORY]
S10-7-G9	729 bp	40	242 aa	29	3p 1d 2m	V F M S	> Hypothetical protein (FL) > Xenopus EST [CDS CATEGORY]
S10-7-H1	347 bp	6	115 aa	5	2p 1d 2m	V F M S	> G2/MITOTIC-SPECIFIC cyclin B1 (FL) [CDS CATEGORY]
S10-7-H10	748 bp	4	249 aa	4	1p 1d 2m	V F M S	> t-box protein like (partial) [CDS CATEGORY]
S10-7-H11	711 bp	6	236 aa	5	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-7-H12	757 bp	4	252 aa	5	3p 1d 2m	V F M S	> Xenopus EST > Marks Related protein (FL) [CDS CATEGORY]
S10-7-H2	724 bp	7	241 aa	7	2p 1d 2m	V F M S	> Zn finger protein (partial) [CDS CATEGORY]
S10-7-H3	732 bp	10	243 aa	7	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-7-H4	728 bp	2	242 aa	3	2p 1d 2m	V F M S	> B99 protein (FL) [CDS CATEGORY]
S10-7-H5	730 bp	7	243 aa	6	3p 1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-7-H6	698 bp	1	232 aa	1	2p 1d 2m	V F M S	> fatty aldehyde dehydrogenase (FL) [CDS CATEGORY]
S10-7-H7	721 bp	8	240 aa	9	1p 1d 2m	V F M S	> Zn finger protein (partial) [CDS CATEGORY]
S10-7-H8	710 bp	1	236 aa	2	1p 1d 2m	V F M S	> methionine aminopeptidase (partial) [CDS CATEGORY]
S10-7-H9	763 bp	12	254 aa	11	1p 1d 2m	V F M S	> leucine-rich acidic nuclear protein (FL) > Duplicate [CDS

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T02240" 2450T650

S10-8-B12	670 bp	11	223 aa	9	1p 1d 2m	V F M S	> delta/YY1 Zn finger protein (FL) [CDS CATEGORY]
S10-8-B2	699 bp	2	232 aa	2	1d 2m	V F M S	> xenopus EST > Duplicate [CDS CATEGORY]
S10-8-B3	694 bp	9	231 aa	9	1d 2m	V F M S	> RAR gamma B (partial)
S10-8-B4	666 bp	3	221 aa	3	2p 1d 2m	V F M S	> RNA binding protein mARK (partial) [CDS CATEGORY]
S10-8-B5	668 bp	3	222 aa	3	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-8-B6	736 bp	2	245 aa	3	1d 2m	V F M S	> Xenopus EST
S10-8-B7	730 bp	5	243 aa	4	1d 2m	V F M S	> Ribosomal protein S6 (FL)
S10-8-B8	721 bp	8	240 aa	8	1p 1d 2m	V F M S	> EIF 4 gamma 2 (FL) [CDS CATEGORY]
S10-8-B9	675 bp	1	224 aa	1	1p 1d 2m	V F M S	> sorting nexin 14 (partial) [CDS CATEGORY]
S10-8-C1	543 bp	2	180 aa	1	1d 2m	V F M S	> Xenopus EST
S10-8-C10	691 bp	76	230 aa	62	2p 1d 2m	V F M S	> delta tubulin (FL) > Xenopus EST [CDS CATEGORY]
S10-8-C11	663 bp	1	220 aa		3p 1d 2m	V F M S	> KIAA0592 related (partial) [CDS CATEGORY]
S10-8-C12	667 bp	4	222 aa	4	1p 1d 2m	V F M S	> pre-mrna splicing factor [CDS CATEGORY]
S10-8-C2	708 bp	10	235 aa	7	1p 1d 2m	V F M S	> hnRNP A1 (FL) [CDS CATEGORY]
S10-8-C3	688 bp		229 aa	1	1p 1d 2m	V F M S	> xenopus ORF > retrotransposon [CDS CATEGORY]
S10-8-C4	690 bp	105	229 aa	73	1d 2m	V F M S	> Xenopus EST
S10-8-C5	686 bp	1	228 aa	2	1p 1d 2m	V F M S	> glucosylphosphotransferase (partial) [CDS CATEGORY]
S10-8-C6	739 bp	3	246 aa	4	3p 1d 2m	V F M S	> Hypothetical ORF (FL) [CDS CATEGORY]
S10-8-C7	649 bp		216 aa	1	2p 1d 2m	V F M S	> cgi-74 like (FL) > Xenopus EST [CDS CATEGORY]
S10-8-C8	726 bp	6	241 aa	3	2p 1d 2m	V F M S	> k1aa0092 related (FL) [CDS CATEGORY]
S10-8-C9	673 bp	3	224 aa	3	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-8-D1	669 bp	2	222 aa	2	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-8-D10	674 bp	2	224 aa	1	1p 1d 2m	V F M S	> beta-catenin (FL) [CDS CATEGORY]
S10-8-D11	619 bp		206 aa	1	3p 1d 2m	V F M S	> Xenopus EST > DMPK-like (partial) [CDS CATEGORY]
S10-8-D12	660 bp	2	219 aa	2	1d	V F M S	> Xenopus EST [CDS CATEGORY]
S10-8-D2	700 bp		233 aa	1	1p 1d 2m	V F M S	> adp-ribosylation factor 6 (FL) [CDS CATEGORY]
S10-8-D3	692 bp	1	230 aa	1	1p 1d 2m	V F M S	> k1aa0948 like (partial) > Xenopus EST [CDS CATEGORY]
S10-8-D4	687 bp		228 aa		1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-8-D5	735 bp	3	244 aa	2	1p 1d 2m	V F M S	> SNF5 protein (FL) [CDS CATEGORY]
S10-8-D6	659 bp	6	219 aa	6	1d 2m	V F M S	> xenopus EST > Conserved Element [CDS CATEGORY]



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S10-8-D7	729 bp	3	242 aa	3	2p 1d 2m	V F M S
S10-8-D8	312 bp	3	103 aa	2	1d 2m	V F M S
S10-8-D9	210 bp		69 aa		1d 2m	V F M S
S10-8-E1	686 bp	4	228 aa	4	1p 1d 2m	V F M S
S10-8-E10	651 bp	2	216 aa	2	2p 1d 2m	V F M S
S10-8-E11	650 bp	8	216 aa	7	1p 1d 2m	V F M S
S10-8-E12	662 bp		220 aa	1	1p 1d 2m	V F M S
S10-8-E2	696 bp	15	231 aa	13	1d 2m	V F M S
S10-8-E3	692 bp	3	230 aa	3	1p 1d 2m	V F M S
S10-8-E4	678 bp	5	225 aa	4	1p 1d 2m	V F M S
S10-8-E5	737 bp	4	245 aa	3	1d 2m	V F M S
S10-8-E6	689 bp	89	229 aa	70	1d 2m	V F M S
S10-8-E7	671 bp	2	223 aa	3	2p 1d 2m	V F M S
S10-8-E8	670 bp		223 aa	1	1p 1d 2m	V F M S
S10-8-E9	673 bp	1	224 aa	2	3d	V F M S
S10-8-F1	696 bp	1	231 aa	1	1p 1d 2m	V F M S
S10-8-F10	668 bp		222 aa	1	1p 1d 2m	V F M S
S10-8-F11	661 bp		220 aa	1	1d 2m	V F M S
S10-8-F12	651 bp	4	216 aa	3	3p 1d 2m	V F M S
S10-8-F2	678 bp	4	225 aa	4	1d 2m	V F M S
S10-8-F3	693 bp	3	230 aa	3	1d 2m	V F M S
S10-8-F4	696 bp	5	231 aa	5	1p 1d 2m	V F M S
S10-8-F5	679 bp	1	226 aa	2	2p 1d 2m	V F M S
S10-8-F6	689 bp	72	229 aa	54	3p 1d 2m	V F M S
S10-8-F7	668 bp	3	222 aa	3	1d 2m	V F M S
S10-8-F8	691 bp	148	230 aa	99	1d 2m	V F M S
S10-8-F9	672 bp	2	223 aa	2	1p 1d 2m	V F M S
S10-8-G1	682 bp	1	227 aa	2	1d 2m	V F M S
S10-8-G10	671 bp	1	223 aa	2	1d 2m	V F M S

> xenopus EST > hypothetical ORF [CDS CATEGORY]
> cdc47-2 (partial) [CDS CATEGORY]
> Xenopus EST
> ASF-2 (FL) [CDS CATEGORY]
> kiaa0995 like (FL) [CDS CATEGORY]
> RAD54 (FL) [CDS CATEGORY]
> Exostosin-2 like (nFL) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> pecanex 1 (partial) [CDS CATEGORY]
> hFetal Brain ORF (nFL)) [CDS CATEGORY]
> Xenopus EST
> Xenopus EST
> Xenopus EST > EMP [CDS CATEGORY]
> human autoantigen like (FL) [CDS CATEGORY]
> Xenopus EST
> ubiquitin-conjugating enzyme e2 (FL) [CDS CATEGORY]
> PKC inhibitor 1 (FL) [CDS CATEGORY]
> Xenopus EST
> Xenopus EST > arginine methyltransferase (partial) [CDS C
> Xenopus EST [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> antizyme inhibitor (nFL) [CDS CATEGORY]
> Nucleosome assembly factor related (FL) [CDS CATEGORY]
> xenopus EST > BRM/BRG related (partial) [CDS CATEGORY]
> Xenopus EST [CDS CATEGORY]
> Xenopus EST
> hnRNP G (FL) [CDS CATEGORY]
> Xenopus EST > Conserved element [CDS CATEGORY]
> Xenopus EST

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S10-8-G11	664 bp	221 aa	1	1p 1d 2m	V F M S	> Fused toes (FL) [CDS CATEGORY]
S10-8-G12	667 bp	222 aa	3	1d 2m	V F M S	> Xenopus EST
S10-8-G2	694 bp	231 aa	1	1d	V F M S	> Xenopus EST [CDS CATEGORY]
S10-8-G3	695 bp	231 aa	2	1p 1d 2m	V F M S	> thymic dendritic cell-derived factor 1 (FL) [CDS CATEGORY]
S10-8-G4	692 bp	230 aa	4	1p 1d	V F M S	> AKT2 Kinase (partial) [CDS CATEGORY]
S10-8-G5	700 bp	233 aa	61	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-8-G6	735 bp	244 aa	3	1p 1d 2m	V F M S	> histone binding protein N1/N2 (FL) [CDS CATEGORY]
S10-8-G7	724 bp	241 aa	3	1d 2m	V F M S	> conserved element > Xenopus EST [CDS CATEGORY]
S10-8-G8	671 bp	223 aa	4	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-8-G9	667 bp	222 aa	3	1p 1d 2m	V F M S	> Cyt C oxidase chain 1 (FL) [CDS CATEGORY]
S10-8-H1	696 bp	231 aa	5	1d 2m	V F M S	> Xenopus EST
S10-8-H10	626 bp	208 aa	4	1p 1d 2m	V F M S	> cytochrome c oxidase subunit II (nFL) [CDS CATEGORY]
S10-8-H11	662 bp	220 aa	2	1d 2m	V F M S	> xenopus EST > Conserved Element [CDS CATEGORY]
S10-8-H12	665 bp	221 aa	2	1p 1d 2m	V F M S	> LIM domain protein (FL) [CDS CATEGORY]
S10-8-H2	697 bp	232 aa	5	1d 2m	V F M S	> Xenopus EST [CDS CATEGORY]
S10-8-H3	693 bp	230 aa	3	1p 1d 2m	V F M S	> phosphoprotein phosphatase 2A regulatory subunit (FL) [CD
S10-8-H4	688 bp	229 aa	4	1p 1d 2m	V F M S	> xETS-2A-2 (FL) [CDS CATEGORY]
S10-8-H5	729 bp	242 aa	3	1p 1d 2m	V F M S	> ras Activating protein (FL) [CDS CATEGORY]
S10-8-H6	728 bp	242 aa	5	1d 2m	V F M S	> Xenopus EST
S10-8-H7	674 bp	224 aa	2	2p 1d 2m	V F M S	> Hypothetical C elegans (Partial) [CDS CATEGORY]
S10-8-H8	737 bp	245 aa	93	1d 2m	V F M S	> Xenopus EST
S10-8-H9	673 bp	224 aa	8	2p 1d 2m	V F M S	> prp28, U5 snrnp (partial) [CDS CATEGORY]

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## APPENDIX 2

Tables S10-1 to S10-8; each table having 4 pages (page 5 of 5 omitted in all cases)

094043.0001  
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**609**

Q9925904.seq

TTTTTTGAAANCCNTNTACNTGTTCCNTTTGCAGGATCCCATCGATTTCGAATTCGATCGCCCCACGCCGTCGGGGGAAAAAGGAGTGT  
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GACGCGGAGCCGGGGACACCGAGCCGAGCGGCTTGTAATCCACAACCTGGTTCTTTGGAAACTGTTGAAAAATGGCAACTCAA  
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GTATTCATTTCTGGAGCAACCAACACAGCAACCATCTTTGAGTGGCAAGGAAATCCAGAGGATGATGATGTGGATCCCAACCAAAT  
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